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October 21, 2002, 09:47:36; Search time 25 Seconds (without alignments) 207.594 Million cell updates/sec
                                                                                                                                                                                                                                                                                                                                                                               562222
GenCore version 5.1.3
Copyright (c) 1993 - 2002 Compugen Ltd.
                                                                                                                                                                                                 US-09-720-828A-4
147
1 APTSSSTKKTQLQLEHLLLDLQMILNGINN 30
                                                                                                                                                                                                                                                                                                                                                                               Total number of hits satisfying chosen parameters:
                                                                                                                                                                                                                                                                                                                                      562222 seqs, 172994929 residues
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
                                                                               OM protein - protein search, using sw model
                                                                                                                                                                                                                                                                                  BLOSUM62
Gapop 10.0 , Gapext 0.5
                                                                                                                                                                                                                                                                                                                                                                                                                       Minimum DB seq length: 0
Maximum DB seq length: 200000000
                                                                                                                                                                                                   Title:
Perfect score:
Sequence:
                                                                                                                                                                                                                                                                                    Scoring table:
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                                                                                                                            Run on:
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sp_mnumeat::
 sp_mnc:*
 sp_organelle:*
 sp_phage:*
 sp_plant:*
 sp_vodent:*
 sp_volunts:*
 sp_vortebrate:*
 sp_vorteb SPTREMBL_19:*

1: sp_archea:*
2: sp_bacteria:*
3: sp_fungi:*
4: sp_human:*
5: sp_invertebrate:*
6: sp_mammal:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

		Description		Q9xs38 papio hamad	_		Q9xt83 halichoerus	Q9bg74 canis famil	Q9tv12 canis famil	Q9mzr9 oryctolagus	Q923t2 sigmodon hi	070329 mesocricetu	Q9ucf5 homo sapien	Q9xt84 delphinapte	088210 cavia porce	Q9gjr4 ovis aries	Q9g183 capra hircu	Q95mp4 ovis aries
SUMMARIES		ID	Q9C001	Q9XS38	Q13169	016334	Q9XT83	Q9BG74	Q9TV12	Q9MZR9	Q923T2	070329	Q9UCF5	O9XT84	088210	Q9GJR4	Q9GL83	Q95MP4
		DB	4	9	4	4	9	9	9	9	11	11	4	9	11	9	9	9
		Query Match Length DB	150	154	156	139	155	99	79	133	155	138	23	154	152	69	155	155
	æ	Query Match	100.0	98.6	92.2	91.2	73.8	73.1	72.8	72.1	72.1	70.1	61.9	56.5	55.8	53.7	53.7	53.7
		Score	147	145	135.5	134	108.5	107.5	107	106	106	103	91	83	82	79	79	79
		Result No.		7	m	4	S	9	7	ω	6	10	11	12	13	14	15	16

RESULT 2

Q95kp3 bubalus bub Q9bq73 canis famil P70291 mus musculu Q9qus8 mus musculu P70294 mus musculu P70293 mus musculu P70293 mus musculu P70293 mus musculu Q9138 trichechus Q9138 trichechus Q9165 strongyloce P71599 mycobacteri O60244 homo sapien Q9unb3 caenorhabdi Q10417 caenorhabdi Q20417 caenorhabdi Q20417 caenorhabdi Q20417 caenorhabdi Q20417 caenorhabdi Q20417 caenorhabdi Q9xsf vibrio chol Q95ky subrio chol Q95ky subrio chol Q96yx subrio chol Q96yx subrio chol Q90yt6 staphylococ Q9uu99 schizosacch Q9vu76 drosophila	cRc64; Length 150; Tindels 0; Gaps 0;
095KP3 098G73 P70291 090USB P70293 P70293 P70293 0910K5 0930K5 0930K5 0930K5 0910M2 091DM2 091DM2 091DM2 091DM2 091DM2 091DM2 091DM3 091DM3 091DM3 092BV0 022042 0978K5 096192 0978K5 096192 0978K5 096193 096199	PRT; 150 AA. Created) Last sequence update) Last annotation and Last annotation Last
53.7 155 6 46.9 150 11 46.9 150 11 45.6 150 11 42.5 159 11 42.5 159 11 35.4 737 16 34.7 30 16 34.0 145 4 34.0 1454 4 34.0 1454 4 33.3 458 5 33.3 543 5 33.3 1130 5 32.7 441 16 32.7 441 16 32.0 39 16 32.0 100 3 32.0 163 3 32.0 163 3 32.0 163 3 32.0 163 3 32.0 163 3 32.0 163 3	PRELIMINARY; (TrEMBLrel. 17 (TrEMBLrel. 17 (TrEMBLrel. 19 2 (FRAGNERY). 2 (FRAGNERY). 2 (FRAGNERY). 3 (FRAGNERY). 4 (Human). 606; 606; 606; 606; 606; 7 NA. 606; 606; 606; 606; 606; 606; 606; 606
117 79 118 669 220 669 221 662.5 23 662.5 24 60.5 25 26 72 26 23 30 30 30 50 31 860.5 24 89 35 449 48 48 36 47 47 48	RESULT 1 09C001 ID 09C001 DT 01-UNN-2001 (TIEMBL DT 01-UNN-2001 (TIEMBL DT 01-DEC-2001 (TIEMBL DE INTERLEUKIN-2 (FRAG C MARMMALIAL ENCHORIA) RN MALEZOR; ON NCBL TAXID-9606; RN [1] TAXID-9606; RN [1] TAXID-9606; RN MALESANZ F., Delgad RT MALLelic Selection RT MALLelic Selection RT MALLelic Selection DR MENS; PO1585; 31NK. DR HSSP; PO1585; 31NK. DR PRINTS; PR00265; IL2; DR PRINTS; PR00265; IL2; DR PRINTS; PR00189; IL2; DR
	# 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0

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92.2%;
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                                                                                               Query Match
Best Local Similarity 90.97
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local Similarity 93.33
Matches 28; Conservative
                                                                                                                                                                                                                                                                                                                                       PRELIMINARY;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Homo sapiens (Human)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQUENCE FROM N.A.
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                                                                                                                                                                                                                                                                                                                                     Q16334
Q16334;
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Q9XT83
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                                                                                                                                                                                                                                                                                                                                          Murillo L.A., Hernandez E., Echeverry S.J., Mendez J.A., Moreno A., Batarroyo M.E.; Patarroyo M.E.; Beheverry S.J., Mendez J.A., Moreno A., Submitted (FBB-1997) to the EMBL/GenBank/DDBJ databases.
EMBL; U88365; AAD41538.1; --
EMBL; U88361; AAD41534.1; --
EMBL; U88363; AAD41536.1; --
EMBL; U88363; AAD41536.1; --
EMBL; U88363; AAD41537.1; --
                                                                                                                                                        Actus lemurinus (Northern gray-necked night monkey),
Actus nancymaae (Owl monkey),
Actus nancymaae (Owl monkey),
Actus nancymaae (Dlack-headed night monkey), and
Actus vociferans (noisy night monkey),
Eukaryota, Metazoa; Chordata, Craniata, Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Cercopithecidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Homo sapiens (Human).
Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi,
Mammalia, Eutheria, Primates, Catarrhini, Hominidae, Homo.
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Submitted (APR-1995) to the EMBL/GenBank/DDBJ databases.
EMBL; U25676; AAA70092.1; -.
HSSP; P01585; 31NK.
InterPro; IPR000779; Interleukin-2.
Pfam; PP00715; IL2; 1.
PRINTS; PR00265; INTERLEUKIN2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PROSITE; PS00424; INTERLEUKIN_2; 1.
SEQUENCE 154 AA; 17675 MW; AB752ABBADA96469 CRC64;
                                                      01-NOV-1999 (TrEMBLrel. 12, Created)
01-NOV-1999 (TrEMBLrel. 12, Last sequence update)
01-OCT-2001 (TrEMBLrel. 18, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Created)
Last sequence update)
Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      98.6%; Score 145; DB 6; L 96.7%; Pred. No. 2.9e-14; live 1; Mismatches 0;
                       154 AA.
                                                                                                                                                                                                                                                                                     Cercopithecinae; Papio.
NCBL_TaxID-9557, 43147, 37293, 57175, 57176;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        156 AA
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                       PRT;
                                                                                                                                        Papio hamadryas (Hamadryas baboon),
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          InterPro; IPR000779; Interleukin-2. Pfam; PF00715; IL2; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PRINTS; PR00265; INTERLEUKIN2.
ProDom; PD003649; Interleukin-2; 1.
SMART; SM00189; IL2; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRT;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Local Similarity 96.7 hes 29; Conservative
                     PRELIMINARY;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRELIMINARY;
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                                         09xs38;
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                   09xs38
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09XS38
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"Molecular cloning and phylogenetic analysis of beluga whale
(Delphinapterus leucas) and grey seal (Halichoerus grypus) interleukin 2.";
                                                                                                                                                                                                                                                                                                                                                              Gaps
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*** Bizenberg O., Faber-Elman A., Lotan M., Schwartz M.;

*** Bizenberg O., Faber-Elman A., Lotan M., Schwartz M.;

*** Thicorleukin-2 transcripts in human and rodent brains: possible captession by astrocytes.";

*** Thicorleukin-2 transcripts in human and rodent brains: possible to a variochem 64:1928-1936(1995).

*** Representation by astrocytes.";

*** Resp. Pol365; 31NK.

*** Representation by astrocytes.";

*** Representation by astrocytes.";

*** Representation by astrocytes.";

*** PROPORTS: PRO0265; INTERLEUKIN2.**

*** PROMOTH PRO03649; Interleukin-2; 1.**

*** RANGH SAMART; SMO189; IL2; 1.**

*** PROMOTHS: PRO0424; INTERLEUKIN_2; 1.**

*** PROMOTHS: P
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Mammalia; Butheria; Primates; Catarrhini; Hominidae; Homo.
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Eukaryota, Metazoa, Chordata, Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Carnivora; Pinnipedia; Phocidae; Halichoerus.
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                                                                                                                                                                                                                                                             DB 4; Length 156;
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Pred. No. 1.2e-12;
0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                         Indels
Prodom; PD003649; Interleukin-2; 1.
SMART; SM00189; IL2; 1.
PROSITE; PS00424; INTERLEUKIN_2; 1.
SEQUENCE 156 AA; 18002 MW; 8E0452D43B336389 CRC64;
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SEQUENCE 139 AA; 15986 MW; 731FBA406D0C63C5 CRC64;
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Last annotation update)
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01-NOV-1999 (TrEMBLrel. 12, Last sequence update)
01-DEC-2001 (TrEMBLrel. 19, Last annotation update)
INTERLEGIKIN 2.
                                                                                                                                                                                                                Score 135.5; DB 4;
Pred. No. 8.1e-13;
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                                                                                                                                                                                                                                                                                                                                                                                                                                          1 APTSSS---TKKTQLQLEHLLLDLQMILNGINN 30
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Gaps

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us-09-720-828a-4.rspt

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SEQUENCE FROM N.A.

PISSUE-SPLEEN, LYMH NODE;

PERVINS 4.2034414; PubMed=10843729;

Perkins H.D., van Leeuwen B.H., Hardy C.M., Kerr P.J.;

Perkins H.D., van Leeuwen B.H., Hardy C.M., Kerr P.J.;

The complete CDNs Sequences of IL-2, IL-4, IL-6 and IL-10 from the European rabbit (Oryctolagus cuniculus).";

Cytokine 12:555-565(2000).

EMBL; AP169168; ARF86552.1; -.

EMSP; PO1585; 31AR.

InterPro; IRR000779; Interleukin-2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                01-DEC-2001 (TrEMBLrel. 19, Created)
01-DEC-2001 (TrEMBLrel. 19, Last sequence update)
01-DEC-2001 (TrEMBLrel. 19, Last annotation update)
INTERLEUKIN 2.
Signodon hispidus (hispid cotton rat).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Sigmodontinae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Oryctolagus cuniculus (Rabbit).
Eukaryota, Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Lagomorpha; Leporidae; Oryctolagus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length 133;
                                                                                                                                                                                                                                  Score 107; DB 6; Length 79;
Pred. No. 9e-09;
6; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 3: Indels
"Cloning and sequencing of canine IL-2.";
Submitted (SEP-1998) to the EMBL/GenBank/DDBJ databases.
EMBL; AF09131; AAD46989.1; -.
HSSP; PO1585; 11RL.
III.erFro: 1PR00779; Interleukin-2.
Pfam; PF00715; IL2; 1.
PRINTS; PR00265; INTERLEUKIN2.
ProDom; PD003649; Interleukin-2; 1.
SMART; SMO0189; IL2: 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Prodom; PF00715; IL2; 1.
Prodom; PD003649; Interleukin-2; 1.
SWART; SM00189; IL2; 1.
SROGHZE; PS00424; INTERLEUKIN_2; 1.
SEQUENCE 133 AA; 14748 MW; 0054758C190B5655 CRC64;
                                                                                                                                                                     NON_TER 1 1
NON_TER 79 79 79
SEQUENCE 79 AA; 9087 MW; 83079BF8F8A659BD CRC64;
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01-OCT-2000 (TrEMBLrel. 15, Last sequence update)
01-DEC-2001 (TrEMBLrel. 19, Last annotation update)
INTERLEUKIN 2 VARIANT IL2DELTA2.
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                                                                                                                                                                                                                                                                                                                                                                                                                     133 AA.
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                                                                                                                                                                                                                                                                                                    3 TSSSTKKTQLQLEHLLLDLQMILNGINN 30
                                                                                                                                                                                                                                                                                                                      PRT;
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Local Similarity 71.4%;
hes 20; Conservative
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Matches
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Q923T2
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"Expression of canine interleukin-2 mRNA in concanavalin A-stimulated canine largession of canine interleukin-2 mRNA in concanavalin A-stimulated canine lamphocytes.",
submitted (JAN-2001) to the EMBL/GenBank/DDBJ databases.
EMBL, AFSAIJIT, AAKO1437.1; -.
HSSP: P01585: IRL.
InterPro; IPR00779; Interleukin-2.
PRINTS; PR000265; INTERLEUKIN2.
SPRAFT, SM00189; IL2; 1.
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01-MAY-2000 (TrEMBLrel. 13, Last sequence update)
01-MAY-2010 (TrEMBLrel. 18, Last sequence update)
01-CT-2001 (TrEMBLRel. 18, Last annotation update)
INPERLEGKIN-2 (FRAGMENT).
Canis familiaris (Dog).
ENKaryota; Metazoa: Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Euterla; Carnivora; Fissipedia; Canidae; Canis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Carnivora; Fissipedia; Canidae; Canis. NCBI_TaxID=9615;
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Pred. No. 6.3e-09;
6; Mismatches 2; Indels :
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SEQUENCE FROM N.A.
German A.J., Helps C.R., Harley R., Hall E.J., Day M.J.;
Vet. Immunol. Immunopathol. 67:385-394(1999).
EMBLA AF072871. AAD40848.1; -.
EMBLA AF072871. AAD40848.1; -.
InterPro. IPR0000779; Interleukin-2.
Prom. PF00715; ILC: IL.
PRINTS; PR00265; ILC: IL.
PRODOM; PD003649; ILC: IL.
PROSTRE: SA001849; ILC: ILRELEUKIN-2; I.
SWART; SMO0189; ILC: INTERLEUKIN-2; I.
SROGTRE: PS00424; IRVERIEUKIN-2; I.
SEQUENCE 155 AA; 17860 MM; F18F449AC672241A CRC64;
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NON_TER 66 66 66 522A893F79DA2AE47 CRC64;
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01-JUN-2001 (TrEMBLrel. 17, Created)
01-JUN-2001 (TrEMBLrel. 17, Last sequence update)
01-DEC-2001 (TrEMBLrel. 19, Last annotation update)
INTERLEUKIN 2 (FRAGMENT).
                                                                                                                                                                                                                                                                                                                                                                     66 AA.
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                                                                                                                                                                                                                                                                     21 APTTSSSTKETQQQLEQLLLDLRLLLLNGVNN 51
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Best Local Similarity 71.0%;
Matches 22; Conservative
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TISSUE-BLOOD;
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SEQUENCE FROM N.A.
MEDILINE-99221046; PubMed-10206205;
St.Laurent G., Beliveau C., Archambault D.;
St.Laurent G., Beliveau G., Archambault D.;
(Molecular cloning and phylogenetic analysis of beluga whale (Delphinapterus leucas) and grey seal (Halichoerus grypus) interleukin
                                                                                                                                                                                                                                                                Gaps
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088210;
01-NOV-1998 (TrEMBLrel. 08, Created)
01-NOV-1998 (TrEMBLrel. 08, Last sequence update)
01-OCT-2001 (TrEMBLrel. 18, Last annotation update)
01-OCT-2001 (TrEMBLrel.)
01-OCT-200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Cetartiodactyla; Cetacea; Odontoceti;
Monodontidae; Delphinapterus.
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                                                             MEDLINE-92289963; PubMed-8512072;
Mullner S., Karbe-Thonges B., Tripier D.;
Mullner S., Karbe-Thonges B., Tripier D.;
Scharge heterogeneity of insulin fusion proteins expressed is Escherichia coli is not due to proteolytic degradation.";
Anal. Biochem. 210.366-373(1993).
SEQUENCE 23 AA; 2637 MW; 40B64C6875CE021F CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Length 154;
                                                                                                                                                                                                   61.9%; Score 91; DB 4; Length 23; 90.5%; Pred; No. 7.1e-07; Mismatches 2; Indels
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ProDom; PD003649; Interleukin-2; 1.
SMART; SM00189; ILZ; 1.
PROSTIE; PS00424; INTERLEUKIN_2; 1.
SEQUENCE 154 AA; 17652 MW; 4288D3D41D04F172 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         01-NOV-1999 (TrEMBLrel. 12, Created)
01-NOV-1999 (TrEMBLrel. 12, Last sequence update)
01-DEC-2001 (TrEMBLrel. 19, Last annotation update)
INTERLEMENT 2.
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ilarity 56.7%; Pred. No. 7.9e-05;
Conservative 6; Mismatches 7;
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EMBL, AF072870; AAD40847.1;
HSSP, POLS65, 31NK.
InterPro; IPR000779; Interleukin-2.
Pfam; PF00715; IL2; 1.
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                                                                                                                                                                                                                                                                                                         3 TSSSTKKTQLQLEHLLLDLQM 23
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Matches 17; Conserv
                                                                                                                                                                                                                                         Best Local Similarity
Matches 19; Conserv
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SEQUENCE FROM N.A.
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    NCBI_TaxID-9606;
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                                               SEQUENCE.
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                                                           SEQUENCE FROM N.A.

Darnell M.R., Pletneva L.M., Langley R.J., Blanco J.C., Prince G.A.;
"Clooning, expression and purification of cotton rat IL-2.";
Submitted (JUL-2001) to the EMBL/GenBank/DDBJ databases.

EMBL; AF398549; AAK94012.1; -.
SEQUENCE 155 AA, 17627 MW; ACADEA865E993291 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MEDLINE-98234044; PubMed-9573100;
Melby P.C., Tryon V.V., Chandrasekar B., Freeman G.L.;
Melby P.C., Tryon V.V., Chandrasekar B., Freeman G.L.;
"Cloning of Syrian hamster (Mesocricetus auratus) cytckine cDNAs and
analysis of cytckine mRNA expression in experimental visceral
leishmaniasis.":
Infect. Immun. 66:2135-2142(1998).
EMBL; AF046212; AAC40097.1; -.
InterPro; IPR000779; Interleukin-2.
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01-A0G-1998 (TrEMBLrel. 07, Last sequence update)
01-A0G-1998 (TrEMBLrel. 18, Last annotation update)
INTERLEUKIN-2 (FRAGMENT)

BASOCILCOCHUS auratus (Golden hamster).

Bukaryota; Metazoos; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Cricetinae;
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01-MAY 2000 (TrEMBLrel. 13, Last sequence update)
01-UMY-2000 (TrEMBLrel. 14, Last annotation update)
INTERLEUKIN 2 (FRAGMENT).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Pred. No. 6.4e-08;
2; Mismatches 6; Indels
                                                                                                                                                                                                                   72.1%; Score 106; DB 11; Length 155; larity 73.3%; Pred. No. 2.5e-08; Conservative 3; Mismatches 5; Indels
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PRINTS; PR00205; INTERLEUKIN2.
PLODOm; PD003649; Interleukin-2; 1.
SMART; SM0189; IL2; 1.
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PROSITE; PSO0424; INTERLEUKIN_2; 1.
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Local Similarity 73.3%;
hes 22; Conservative
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es 22; Conserv
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Sigmodon.
NCBI_TaxID-42415;
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SEQUENCE
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Q9UCF5
ID Q9UCF5
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Matches
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070329
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Search completed: October 21, 2002, 09:49:35 Job time : 27 secs
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Luehken G., Prinzenberg E.-M., Hiendleder S., Erhardt G.;
"A single strand conformation polymorphism in the ovine interleukin;
(IL-2) gene.";
J. Anim. Sci. 0.0-0(2000).
EMBL; AF213687; AAG43986.1; -.
EMBL; AF213883; AAG35709.1; -.
HSSP; PO1589; IRL.
Interpro; IPR00079; Interleukin-2.
PRINTS; PR000265; INTERLEUKIN2.
SPINONS; PR000869; INTERLEUKIN2.
SMART; SM00189; IL2; 1.
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Wakaryota: Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Cefartiodactyla; Ruminantia; Pecora; Bovoidea;
Bovidae; Caprinae; Ovis.
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Pred. No. 0.00014;
6; Mismatches 8; Indels

        SIGNAL
        1
        20
        POTENTIAL.

        CHAIN
        21
        152
        INTERLEUKIN 2.

        SEQUENCE
        152 Aa;
        17271 MW;
        CA7ACO8C1B8DDIFA CRC64;

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            01-MAR-2001 (TrEMBLrel. 16, Created)
01-MAR-2001 (TrEMBLrel. 16, Last sequence update)
01-DEC-2001 (TrEMBLrel. 19, Last annotation update)
INTERLEDKIN 2 PRECURSOR (FRAGMENT).
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INTERLEUKIN 2.
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69 AA; 7711 MW; 1
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Best Local Similarity 53.3%;
Matches 16; Conservative
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TISSUE-SPLEEN;
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SEQUENCE
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SIGNAL
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09GL83
ID 09GL8:
AC 09GL8:
DT 01-MA
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A Ying O.H., Li X.R., Pan J.Y.;

Tring O.H., Li X.R., Pan J.Y.;

R Embl., AF3701018; AAC287081.; -.

R Embl., AF3701018; AAC287081.; -.

R HSSP; PO1585; 31NK.

R InterPro; IPR000779; Interleukin-2.

R PAGM: PF00715; ILL2: 1.

R PRINTS: PR00355; INTERLEUKIN-2.

R PRINTS: PR003649; ILL2: 1.

R PRINTS: PR003649; ILL2: 1.

R PRINTS: PR00484; ILL2: 1.

R PROSITE: PS00444; INTERLEUKIN-2; 1.

R PROSITE: PS00444; INTERLEUKIN-2; 1.
                                                                                                      capra hircus (Goat).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Cetartiodactyla; Ruminantia; Pecora; Bovoidea;
Bovidae; Caprinae; Capra.
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01-MAR-2001 (TrEMBLrel. 16, Last sequence update)
01-DEC-2001 (TrEMBLrel. 19, Last annotation update)
INTERLEUKIN 2.
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Matches 16, Conservative
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Sequence 3, Appli Sequence 111, App Patent No. 5314995 Sequence 2, Appli Sequence 6, Appli Sequence 6, Appli Sequence 6, Appli Sequence 18, Appli Sequence 5, Appli Sequence 2, Appli Sequence 2, Appli Sequence 2, Appli

OM protein

Run on:

Sequence:

Searched:

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Sequence 2, Application US/09116594

Patent No. 6168785

GENERAL INFORMATION:
APPLICANT: ECKENBERG, Ralph
APPLICANT: MOREAU, Jean-Louis
APPLICANT: MOREAU, Jean-Claude
TITLE OF INVENTION: BIOLOGICAL APPLICATIONS OF NEW PEPTIDES OF IL-2 AND
TITLE OF INVENTION: DERIVATIVES AND USE AS THERAPEUTIC AGENTS
TITLE OF INVENTION: US/09/116,594
CURRENT APPLICATION NUMBER: US/09/116,594
CURRENT FILING DATE: 1998-07-16
NUMBER OF SEQ ID NOS: 2
SOFTWARE: PATENTIN Ver. 2.1

LENGTH: 31
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 OTHER INFORMATION: Description of Artificial Sequence:peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 13, Application US/08127351

Patent No. 549761

GENERAL INFORMATION:
APPLICANT: BELINKA Jr. BENJAMIN A.
APPLICANT: CUGGHLIN. DANIEL J.
APPLICANT: ALVAREZ, VERNON L.
APPLICANT: WOOD, RICHAND
TITLE OF INVENTION: CONSTRUCTS
NUMBER OF SEQUENCES: 56
CORRESPONDENCE ADDRESS:
ADDRESSEE: DELON, SPIVAK, MCCLELLAND, MAIER &
ADDRESSEE: P.C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Indels
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STREET: 1755 S. Jefferson Davis Highway, Suite 400
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Pred. No. 1.9e-16;
: Mismatches 0;
                                          US-08-818-562-2
US-08-155-888-2
US-07-912-915-2
PCT-US91-05826-2
US-08-875-811-59
US-08-875-811-61
US-08-806-121B-3
US-08-930-917A-18
US-08-930-917A-18
US-08-459-906-5
US-08-170-15-7
US-08-170-15-7
US-08-170-15-7
US-08-170-15-7
US-08-170-15-7
                     US-09-522-217-111
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Matches 30; Conserva
RESULT 2
US-08-127-351-13
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Sequence 15, Appl
Sequence 5, Appl
Sequence 6, Appl
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Patent No. 5210029
Patent No. 5256769
Patent No. 5496934
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(without alignments)
56.367 Million cell updates/sec
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/cgn2_6/ptodata/1/iaa/5B_COMB.pep:*
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       GenCore version 5.1.3
Copyright (c) 1993 - 2002 Compugen Ltd
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1 APTSSSTKKTQLQLEHLLLDLQMILNGINN 30
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Listing first 45 summaries
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5464939-2
5496924-55

    protein search, using sw model

                                                                        October 21, 2002, 09:48:11
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Gapop 10.0 , Gapext 0.5
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Maximum DB seq length: 200000000
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Match Length DB
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                                                                                                                                                                                                                                                            Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: BELINKA JF, BENJAMIN A.
APPLICANT: COUGHLIN, DANIEL J.
APPLICANT: ALVAREZ, VERNON L.
APPLICANT: WOOD, RICHARD.
TITLE OF INVENTION: METAL-BINDING TARGETED POLYPEPTIDE
TITLE OF INVENTION: CONSTRUCTS
NUMBER OF SEQUENCES: 5
CORRESPONDENCE ADDRESS:
ADDRESSEE: NEUSTADY, P.C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            STREET: 1755 S. Jefferson Davis Highway, Suite 400 CITY: Arlington
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CONTINUE TYPE: Floppy disk
MEDIUM TYPE: Floppy disk
MEDIUM TYPE: Floppy disk
COMPUTER: Elba PC compatible
COMPUTER: Elba PC compatible
COMPUTER: DEADLISH: PC-DOS/MS-DOS
SOFTWARE: PATENTIN Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/487,221A
FILING DATE: 07-JUN-1995
CLASSIFICATION: 424
PRIOR APPLICATION NUMBER: US 08/127,351
FILING DATE: 28-SEP-1993
ATTORNEY/ACENT INFORMATION:
NAME: Villacorta, Gilberto M.
REGISTRATION NUMBER: 4980-004-44
TELEPHONE: (703) 413-320
TELEPHONE: CARRATION FRENCHERISTICS
INFORMATION FOR SEQ ID NO: 13:
SEQUENCE CHRRACTERISTICS
ILBOTH: 50 maino acids
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Best Local Similarity 100.0%; Score 147; DB 1;
Best Local Similarity 100.0%; Pred. No. 3.4e-16;
Matches 30; Conservative 0; Mismatches 0;
                                                                                                                                                                                                         Query Match 100.0%; Score 147; DB 1; Best Local Similarity 100.0%; Pred. No. 3.4e-16; Matches 30; Conservative 0; Mismatches 0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                           US-08-487-221A-13; Sequence 13, Application US/08487221A; Patent No. 5593656
                    13:
               INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 50 amino acids
TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TOPOLOGY: unknown
MOLECULE TYPE: peptide
US-08-487-221A-13
                                                                                                                TOPOLOGY: unknown
MOLECULE TYPE: peptide
US-08-480-3678-13
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22202
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                                                                                                                                                                                                                                                                                                                                                                                                                                            RESULT 4
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US-08-480-3678-13
US-08-480-3678-13
Sequence 13, Application US/08480367B
Sequence 13, Application US/08480367B
Setent No. 5578288
Septicant: Condition
APPLICANT: BELINKA Jr, BENJAMIN A.
APPLICANT: ALVAREZ, VERNON L.
APPLICANT: WOOD, RICHARD
TITLE OF INVENTION: METAL-BINDING TARGETED POLYPEPTIDE
TITLE OF INVENTION: CONSTRUCTS
NUMBER OF SEQUENCES: 56
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length 50;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Indels
                                                                                                     COMPUTER READBLE FURKH:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOUTHARE: Patentin Release #1.0, Version #1.25
SURREWI APPLICATION DATE:
APPLICATION NUMBER: US/08/127,351
FILING DATE: 28-SEP-1993
CLASSIFICATION: 534
ATTORNEY/AGENT INFORMATION:
NAME: Villacorta, Gilberto M.
REGISTRATION NUMBER: 4980-004-44
REFERENCE/DOCKET NUMBER: 4980-004-44
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION OF A13-320
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SOFTWARE: Patentin Release #1.0, Version #1.25 CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/480,367B
FILING DATE: 07-06-95
CLASSIFICATION: 424
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local Similarity 100.0%; Pred. No. 3.4e-16; Matches 30; Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADDRESSEE: LOWE, PRICE, LeBLANC & BECKER STREET: 99 Canal Center Plaza, Suite 300 CITY: Alexandria STATE: Virginia COUNTRY: U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FILLING DATE: 07-06-95
CLASSIFICATION: 424
ATTORNEY, AGENT INFORMATION:
NAME: VILLACOTTA, GILDERTO M.
REGISTRATION NUMBER: 34,038
REFERENCE/DOCKET NUMBER: 2654-002A
TELECOMMUNICATION INFORMATION:
TELEPHONE: (703) 684-111
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1 APTSSSTKKTQLQLEHLLLDLQMILNGINN 30
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      : 50 amino acids amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        MOLECULE TYPE: peptide US-08-127-351-13
                                                                                                COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       unknown
CITY: Arlington STATE: Virginia
                         STATE: Virginia COUNTRY: U.S.A. ZIP: 22202
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ZIP: 22314
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ò
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER: IBM 386
COMPUTER: IBM 386
OPERATING SYSTEM: WINDOWS 3.1
SOFTWARE: WORDPERFECT 5.1
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             rrPE: amino acid;
; TOPOLOGY: linear;
; MOLECTUE TYPE: protein
US-08-817-787-15
STATE: New York COUNTRY: USA ZIP: 10020 COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     US-08-160-376A-5
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Sequence 15, Application US/08817787
Sequence 15, Application US/0881787
Sequence 16 Sequence 17 Table OF INVENTION OF TITLE OF INVENTION RECOMBINANT PROTEINS TO MULTI-FUNCTIONAL COMPLEXES CORRESPONDENCE ADDRESS:
Sequence 15, Application US/0881787
Sequence 15,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ö
                                                                                                                       Sequence 13, Application US/08480370

Sequence 13, Application US/08480370

Patent No. 5609847

GENERAL INFORMATION:
APPLICANT: BELIKA Jr, BENJAMIN A. APPLICANT: COUGHLIN, DANIEL J. APPLICANT: ALVAREZ, VERNON L. APPLICANT: MOOD, RICHARD TITLE OF INVENTION: METAL-BINDING TARGETED POLYPEPTIDE TITLE OF INVENTION: CONSTRUCTS NUMBER OF SEQUENCES: 56

CORRESPONDENCE ADDRESS:
ADDRESSEE: OBLON, SPIVAK, MCCLELLAND, MAIER & ADDRESSEE: D.C. STREET: 1755 S. Jefferson Davis Highway, Suite 400

CITY: Alington STATE: Virgina

COUNTRY: U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ; DB 1; Length 50; 3.4e-16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPENTING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/480,370
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FILING WALE:

CLASSIFICATION: 424

PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/127,351

FILING DATE: 28-SEP-1993
ATTORNEY/AGENT INFORMATION:
NAME: VILLACOMINICATION INFORMATION:
TELEPLET ON UNBER: 34,038

REGISTRATION NUMBER: 4980-004-44

TELEPAN: (703) 413-220

TELEFAX: (703) 413-220

TELEFAX: (703) 413-220

TELEFAX: (703) 413-220

TELEFAX: SEQUENCE CHARACTERISTICS:
SEQUENCE CHARACTERISTICS:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           100.0%; Score 147;
ilarity 100.0%; Pred. No. 3
Conservative 0; Mismatche
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1 APTSSSTKKTQLQLEHLLLDLQMILNGINN 30
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amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 unknown
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Best Local Similarity
Matches 30; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ZIP: 22202
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TOPOLOGY:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              RESULT 6
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CURRENT APPLICATION DATA:
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/817,787
FILING DATE: 23-SEP-1997
CLASSIFICATION DATA:
APPLICATION NUMBER: PCT/EP95/04117
FILING DATE: 20-OCT-1994
ATTORN RAPLICATION DATA:
APPLICATION NUMBER: EP 94 11 6558.1
FILING DATE: 20-OCT-1994
ATTORNEY/AGENT INPORMATION:
NAME: HALBY Jr., James F.
REGISTRATION NUMBER: 27,794
REGISTRATION NUMBER: 27,794
REGISTRATION NUMBER: 27,794
RELEPRACE/DOCKET NUMBER: 27,794
RELEPRACE/DOCKET NUMBER: 21,794
RELEPRACE/DOCKET NUMBER: 21,794
RELEPRACE 212-596-9000 MEDIUM TYPE: Floppy disk COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS INFORMATION FOR SEQ ID NO: 15: SEQUENCE CHARACTERISTICS: LENGTH: 88 amino acids

ö Length 88; Indels Query Match 100.0%; Score 147; DB 4; Best Local Similarity 100.0%; Pred. No. 6.8e-16; Matches 30; Conservative 0; Mismatches 0;

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1 APTSSSTKKTQLQLEHLLLDLQMILNGINN 30

3 APTSSSTKKTQLQLEHLLLDLQMILNGINN 32

DISKETTE, 3.5 INCH, 1.44 Mb STORAGE

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GENERAL INFORMATION:
APPLICANT: Ralph, Peter
APPLICANT: Martin, George
APPLICANT: Piatek, Michael
APPLICANT: Larrick, James W.
TITLE OF INVENTION: Multifunctional M-CSF Proteins and Genes Encoding
TITLE OF INVENTION: Therefor
                                                                                                                                                                                                                                                                                                              IMMUNOSTIMULANT AGENT CONTAINING
INTERLEUKIN-2 AND 5'-DEOXY-5-FLUOROURIDINE
                                                                                                                                                                                                         Sequence 1, Application US/07800366
Patent No. 5250296
GENERAL INFORMATION:
GENERAL INFORMATION:
TITLE OF INVENTION: IMMUNOSTIMULANT AGENT CONTAINING
TITLE OF INVENTION: INMUNOSTIMULANT AGENT CONTAINING
TITLE OF INVENTION: INTERLEUKIN-2 AND 5'-DEOXY-5-FLUOROINUMBER OF SEQUENCES: 1
CORRESPONDENCE ADDRESS:
ADDRESSEE: DAVID G, CONLIN; DIKE, BRONSTEIN, ROBERTS
ADDRESSEE: CUGHMAN
STREET: 130 Water Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADDRESSEE: CHIRON CORPORATION
STREET: Intellectual Property - R440, P.O. Box 8097
CITY: Emveryville
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
RAPPLICATION NUMBER: US/07/800,366
FILING DATE: 19911127
CLASSIFICATION: 424
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           100.0%; Score 147; DB 1;
100.0%; Pred. No. 1.1e-15;
Live 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ATTORNEY/AGENT INFORMATION:
NAME: Castle, Donald R
REGISTRATION 124,20
REFERENCE/DOCKET NUMBER: 41417(281)
TELECOMMUNICATION INFORMATION:
TELEFAX: (617)523-3400
TELEFAX: (617)523-6440
TELEFAX: (617)523-6440
TELEFAX: (617)523-6440
TELEFAX: (617)523-6440
TELEFAX: (617)523-6440
TELEFAX: (617)523-6440
TELEFAX: 100531 STRE UR
INFORMATION FOR SEQ ID NO: 1: SEQUENCE CHARACTERISTICS:
LENGTH: 133 amino acids
TYPE: AMINO ACID
                                           1 APTSSSTKKTQLQLEHLLLDLQMILNGINN 30
                                                                    2 APTSSSTKKTQLQLEHLLLDLQMILNGINN 31
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1 APTSSSTKKTQLQLEHLLLDLQMILNGINN 30
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Matches 30; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ; TOPOLOGY: linear
; MOLECULE TYPE: protein
US-07-800-366-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Massachusetts
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CITY: Boston
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ZIP: 02109
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US-08-354-456A-5
                                                                                                                                                             RESULT 9
US-07-800-366-1
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                                                                                                                                                                                                                                                                                                                                                                                                          Length 96;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GENERAL INFORMATION:
APPLICANT: Obermeier, Rainer
APPLICANT: Gerl, Martin
APPLICANT: dedwig, Jurgen
APPLICANT: sabel, Walter
TITLE OF INVENTION: Process for Obtaining Insulin Having
TITLE OF INVENTION: Correctly Linked Cystine Bridges
NUMBER OF SEQUENCES: 12
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                          Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADDRESSEE: Finnegan, Henderson, Farabow, Garrett & ADDRESSEE: Dunner
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  STREET: 1300 I Street, N.W.
CITY: Washington
STATE: 0.C.
COUNTRY: United States of America
ZIP: 20005-3315
COMPUTER READBLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/389,487
FILING DATE:
CLASSIFICATION: 530
                                                                                                                                                                                                                                                                                                                                                                                                     Query Match 100.0%; Score 147; DB 1; Best Local Similarity 100.0%; Pred. No. 7.6e-16; Matches 30; Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Ouery Match

100.0%; Score 147; DB 1;
Best Local Similarity 100.0%; Pred. No. 7.6e-16;
Matches 30; Conservative 0; Mismatches 0;
APPLICATION NUMBER: GE P 4240420.7
FILING DATE: December 2, 1992
ATTORNEY/AGENT INFORMATION:
NAME: BATDARA V MALTEN:
REGISTRATION NUMBER: 31,287
REFERENCE/DOCKET NUMBER: HOE 92/F 384
TELECOMMUNICATION INFORMATION:
TELEPHONE: (908) 231-4079
TELEPHONE: (908) 231-255
INFORMATION FOR SEO ID NO: 5: SEQUENCE CHARACTERISTICS:
LENGTH: 95 Amino Acid (AA)
TYPE: Amino Acid (AA)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1 APTSSSTKKTQLQLEHLILDLQMILNGINN 30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2 APTSSSTKKTQLQLEHLLLDLQMILNGINN 31
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 8, Application US/08389487
Patent No. 5663291
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ATTORNEY/AGENT INFORMATION:
NAME: Einaudi, Carol P.
REGISTRATION NUMBER: 32,220
REFERENCE/DOCKET NUMBER: 0248
TELECOMMUNICATION INFORMATION:
TELEPHONE: 202-408-4400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELEPHONE: 202-408-4000
TELEFAX: 202-408-4400
INFORMATION FOR SEQ ID NO: 8:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             96 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        LENGTH: 96 amino acid
TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ; MOLECULE TYPE: peptide US-08-389-487-8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    linear
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Gaps

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Indels

Length 133;

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Sequence 89, Application US/08318193
Patent No. 5641663
GENERAL INFORMATION:
APPLICANT: MALEK, LAWIENCE T.
TITLE OF INVENTION: OF BIOACTIVE HUMAN GRANULOCYTE MACROPHAGE COLONY
TITLE OF INVENTION: STIMULATING FACTOR (CM-CSF) AND OTHER HETEROLOGOUS
TITLE OF INVENTION: PROTEINS FROM STREPTOMYCES
NUMBER OF SEQUENCES: 91
CORRESPONDANCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER: IBM PC Compatible
COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATCHILIN Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/318,193
FILING DATE:
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/07/935,314
FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local Similarity 100.0%; Pred. No. 1.1e-15;
Matches 30; Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DB 1;
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ER: 18740/116 CACO
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADDRESSEE: Foley & Lardner
STREET: 1800 Diagonal Road, Suite 500
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ATTORNEY/AGENT INFORMATION:
NAME: BENT, STEPHEN A.
RECISTRATION NUMBER: 29,768
REFERENCE/DOCKET NUMBER: 18740/116 CTELEPHONE: (703)683-4109
TELEPHONE: (703)683-4109
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1 APTSSSTKKTQLQLEHLLLDLQMILNGINN 30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1 APTSSSTKRTQLQLEHLLLDLQMILNGINN 30
                                                                                                                                                                                                                                                                                             ; NAME/KEY; Protein
; LOCATION: 1..133
; OTHER INFORMATION: /label= IL2
US-08-225-224-3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       STATE: Virginia
CITY: Alexandria
STATE: Virginia
COUNTRY: USA
ZIP: 22313-029
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
TELEPAONE: (415) 543-9600
TELEPAX: (415) 543-5043
INFORMATION FOR SEQ ID NO: 3:
SEQUENCE CHARACTERISTICS:
LENGTH: 133 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 133 amino acids
amino acid
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                             unknown
                                                                                                                                                     TYPE: amino acid
STRANDEDNESS: unknown
TOPOLOGY: unknown
MOLECULE TYPE: protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              MOLECULE TYPE: protein US-08-318-193-89
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        US-08-318-193-89
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                                                                                                                                                                                                                                                                                   FEATURE:
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; Patent No. 563559
; GENERAL INFORMATION:
    APPLICANT: PASTAN, Ira
APPLICANT: RREITMAN, ROBERT J.
    TITLE OF INVENTION: CIRCULARLY PERMUTATED LIGANDS AND
    TITLE OF INVENTION: CIRCULARLY PERMUTATED FUSION PROTEINS
    NUMBER OF SEQUENCES: 57
    CORRESPONDENCE ADDRESS:
    ADDRESSEE: Townsend and Townsend Khourie and Crew
    STREET: Steuart Street Tower, One Market Plaza
    CONTRY: California
    COUNTRY: US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length 133;
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SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/225,224
FILING DATE: 8-APR-1994
CLASSIFICATION: 530
ATTORNEY/AGENT INFORMATION:
NAME: Weber, Ellen L.
REGISTRATION NUMBER: 32,762
REFEISTRATION NUMBER: 15280-193
TELECOMMUNICATION INFORMATION:
                                    COUNTRY: U.S.A.
21P: 94662-8097
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC Compatible
COMPUTER: ISM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    100.0%; Score 147; DB 1;
100.0%; Pred. No. 1.1e-15;
11ve 0; Mismatches 0;
                                                                                                                                                                                                                                     SUTRENT APPLICATION DATA:
SUTRENT APPLICATION DATA:
APPLICATION DATA:
APPLICATION NUMBER: US/08/354,456A
FILING DATE: 12-DEC-1994
CLASSIFICATION TA:
APPLICATION NUMBER: US 07/995,338
FILING DATE: 21-DEC-1992
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
REGISTRATION NUMBER: 31,395
REFERENCE/DOCKET NUMBER: 750.003/32387
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1 APTSSSTKKTQLQLEHLLLDLQMILNGINN 30
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELEX: n/a
INFORMATION FOR SEQ ID NO: 5
SEQUENCE CHARACTERISTICS:
LENGTH: 133 amino acids
TYPE: amino acid
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Best Local Similarity 100.0
Matches 30; Conservative
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MOLECULE TYPE: protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           US-08-354-456A-5
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ZIP: 94304-1104
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
CMDPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    133 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       LENGTH: 133 amino acids
TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           LENGTH: 133 amino acidd;
TYPE: amino acidd
STRANDEDRESS: single;
TOPOLOGY: linear
MOLECULE TYPE: peptide
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100.0%; Score 147; DB 1; Length 133;
Best Local Similarity 100.0%; Pred. No. 1.1e-15;
Matches 30; Conservative 0; Mismatches 0; Indels 0
                                                   Indels
                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 1, Application US/08284393B
; Sequence 1, 5696234
; GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: ZUTAWSKI, Sandra M.
APPLICANT: ZUTAWSKI, Gerard
TITLE OF INVENTION: MUTEINS OF MAMMALIAN CYTOKINES
NUMBER OF SEQUENCES: 16
CORRESPONDENCE ADDRESS:
ADDRESSEE: DNAX Research Institute
STREET: 901 California Avenue
CITY: Palo Alto
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 2, Application US/08284393B
Patent No. 5696234
GENERAL INFORMATION:
APPLICANT: ZUTAWSKI, Sandra M.
APPLICANT: ZUTAWSKI, Gerard
TITLE OF INVENTION: MUTEINS OF MAMMALIAN CYTOKINES
NUMBER OF SEQUENCES: 16
CORRESPONDENCE ADDRESS:
ADDRESSED: DNAX Research Institute
STREET: SOI California Avenue
CITY: Palo Alto
STATE: California
COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COUNTRY: CALLIOETHORAGE
COUNTRY: USA

ZIP: 94304-1104
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIN PC-DOS/MS-DOS
SOFTWARE: PATENTIN PAPA:
APPLICATION NUMBER: US/08/284,393B
FILING DATE: U1-AUG-1994
CLASSIFICATION: 435
ATORNEY/AGENT INFORMATION:
NAME: CALING ECANID.
REGISTRATION NUMBER: 34,090
REFERENCE/POCKET NUMBER: DX0389
TELEPHONE: 415-82-9196
TELEPHONE: 415-82-9196
    Best Local Similarity 100.0%; Pred. No. 1.1e-15; Matches 30; Conservative 0; Mismatches 0;
                                                                                                                                  1 APTSSSTKKTQLQLEHLLLDLQMILNGINN 30
                                                                                                                                                                                 1 APTSSSTKKTQLQLEHLLLDLQMILNGINN 30
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  single
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      STRANDEDNESS: Signature of the strand of the
                                                                                                                                                                                                                                                                                                                                                          RESULT 13
US-08-284-393B-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             RESULT 14
US-08-284-393B-2
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Sequence 3, Application US/08284393B
Patent NO. 5696234
GENERAL INFORMATION:
APPLICANT: ZULTANSH, Sandra M.
APPLICANT: ZULTANSH, GETALG
TITLE OF INVENTION: MUTEINS OF MAMMALIAN CYTOKINES
NUMBER OF SEQUENCES: 16
CORRESPONDENCE ADDRESS:
ADDRESSEE: DNAX Research Institute
STREET: 901 California Avenue
CITY: Palo Alto
STATE: California
COUNTRY: 0240-1104
COMPUTER: Elabopy disk
COMPUTER: Elabopy disk
COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA: 34,090
FILING DATE: 01-A016-1994
CLASSIFICATION NUMBER: 34,090
FREIERATION NUMBER: 34,090
RESTERRATION NUMBER: 34,090
REFERENCE/DOCKET NUMBER: DX0389
TELECOMMUNICATION INFORMATION:
NAME: Ching, Edwin P.
REJERAX: 415-496-1200
INFORMATION FOR SEQ ID NO: 3:
SEQUENCE CHARACTERIFICS:
LENGTH. 133 aminor action
LENGTH. 133 aminor action
SOFTWARENT APPLICATION DATA:
CURRENT APPLICATION DATA:
APPLICATION UNUMBER:
FILING DATE: 01-804-3938
FILING DATE: 01-804-1994
CLASSIFTCATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: COINING Edwin P.
REGISTRATION NUMBER: 34.090
REGISTRATION NUMBER: 34.090
REGISTRATION NUMBER: 12.000
REGISTRATION NUMBER: 12.000
REGISTRATION NUMBER: 12.000
REGISTRATION NUMBER: 12.000
RELEPHONE: 415-852-9196
TELEPHONE: 415-852-9196
INFORMATION FOR SEQ ID NO: 2: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 147; DB 1;
Pred. No. 1.1e-15;
; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match 100.0%; Score 147; D
Best Local Similarity 100.0%; Pred. No. 1.1
Matches 30; Conservative 0; Mismatches
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; MOLECULE TYPE: peptide US-08-284-393B-3
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0; Gaps Query Match 100.0%; Score 147; DB 1; Length 133; Best Local Similarity 100.0%; Pred. No. 1.1e-15; Matches 30; Conservative 0; Mismatches 0; Indels 0

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Search completed: October 21, 2002, 09:50:15 Job time: 13 secs

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October 21, 2002, 09:47:01; Search time 10 Seconds (without alignments) 116.159 Million cell updates/sec
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GenCore version 5.1.3
Copyright (c) 1993 - 2002 Compugen Ltd.
                                                                                                                                                                                     US-09-720-828A-4
147
1 APTSSSTKKTOLOLEHLLLDLOMILNGINN 30
                                                                                                                                                                                                                                                                                                                                                        Total number of hits satisfying chosen parameters:
                                                                                                                                                                                                                                                                                                                      105224 seqs, 38719550 residues
                                                                                                                                                                                                                                                                                                                                                                                                                                                      Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
                                                                        OM protein - protein search, using sw model
                                                                                                                                                                                                                                                               BLOSUM62
Gapop 10.0 , Gapext 0.5
                                                                                                                                                                                                                                                                                                                                                                                               Minimum DB seq length: 0
Maximum DB seq length: 2000000000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SwissProt_40:*
                                                                                                                                                                                       Title:
Perfect score:
Sequence:
                                                                                                                                                                                                                                                               Scoring table:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Database :
                                                                                                                                                                                                                                                                                                                      Searched:
                                                                                                              Run on:
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

ption	P01585 homo sapien Q29615 macaca fasc P51498 macaca mula P46649 cercocebus o62641 mirounga an Q07885 felis silve Q29416 canis famil O77620 oryctolagus P26891 sus scrofa Q08081 meriones un P26891 meriones un P37997 eguus cabal O97513 orcinus orc P63016 bos taurus P37997 eguus cabal O97513 orcinus orc P63016 bos taurus P19114 ovis aricup P21133 emericella P70859 borrella bu P2133 emericella P70859 borrella bu P27980 simian immu P44581 haemophilus Q92881 chamaydia p Q92mal helicobacte Q92mal helicobacte Q92mal helicobacter P31623 sheep pulmo P87320 schizosacch P98204 arabidopsis P41596 drosophila
SUMMARIES	11.2. HUMAN 11.2. MACFA 11.2. CERTO 11.2. CERTO 11.2. CERTO 11.2. ERECA 11.2. FRAT 11.2. RAT 11.2. P1G 11.2. P1G 11.2. P1G 11.2. P1G 11.2. ANSE 11.2. CAPHI 11.2. SHEEP 11.2. CAPHI 11.2. SHEEP 11.2. CAPHI 11.2. CHEPHI 12. CHEPHI 13. CHEPHI 13. CHEPHI 14. CAPHI 15. CHEPHI 15. CHEPHI 16. CAPHI 16. CAPHI 17. CAPHI 17. CAPHI 18. CAPHI
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ALIGNMENTS

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    SEQUENCE FROM N.A.
SPECIES-Human; TISSUE-placenta;
MEDLINE-96422299; PubMed-8824916;
Chernicky C.L., Tan H., Burfeind P., Ilan J.,
Sequence of interleukin-2 isolated from human placental poly A+ RNA:
possible role in maintenance of fetal allograft.";
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Nishino N., Obaru K., Maeda S., Shimada K., Onoue K.; "Organization of the DNA regions flanking the human interleukin
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Chen S.J., Holbrook N.J., Mitchell K.F., Vallone C.A., Greengard J.S., Crabtree G.R., Lin Y.;
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J. Neurochem. 64:1928-1936(1995).
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MEDLINE-90008901; PubMed-2793860;
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This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation-the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (see http://www.lsb-sib.ch/announce/or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          MEDLINE-9511955; Pubmed-7529123;
Bamborough P., Hedgecock C.J., Richards W.G.;
Bamborough P., Hedgecock C.J., Richards W.G.;
The interleukin-2 and interleukin-4 receptors studied by molecular modelling.";
Tructure 2:839-851(1994).

Structure 2:839-851(1994).

INTOGENIC STIMULATION, THIS PROTEIN IS REQUIRED FOR T-CELL PROCIFERATION AND OTHER ACTIVITIES CRUCIAL TO REGULATION OF THE PROTEIN IS REQUIRED FOR T-CELL PROLIFERATION AND OTHER ACTIVITIES CRUCIAL TO REGULATION OF THE ACTIVATED KILLER CELLS, MANCATES, LYMPHOKINE-CHARD KILLER CELLS, MANCATES, LAWHOKINE-CHARD CELLS.

INMUNE RESPONSE. CAN STIMULATE B CELLS, MONCOYTES, LYMPHOKINE-CHARD CAN STIMULANE B CELLS, MONCOYTES, LYMPHOKINE-CHARD CELLS.

INMUNE RESPONSE. CAN STIMULATE B CELLS, MONCOYTES, LYMPHOKINE-CHARD CELLS.

CHARACTERIZED BY A CHROMOSOMAL TRANSLOCATION T(4,16)(Q26;P13)
WHICH INVOLVES ILZ AND BGWA.

IN PARIANCEUTICAL: Available under the name Proleukin (Chiron). Used to patients with renal cell carcinoma or metastatic melanoma.

INMM-"http://www.rndsystems.com/asp/g_sitebuilder.asp?bodyId=206".
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mott H.R., Driscoll P.C., Boyd J., Cooke R.M., Weir M.P.,
Campbell I.D.;
"Secondary structure of human interleukin 2 from 3D heteronuclear NMR
   qene of a cell
"A viral long terminal repeat in the interleukin 2 line that constitutively produces interleukin 2.";
                                  line that constitutively produces interleukin 2."
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AAA35454.1; -.
AAA59139.1; -.
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MEDLINE-88070646; PubMed-3500515;
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MEDLINE-92335891; PubMed-1631562;
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Biochemistry 31:7741-7744(1992).
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A01849; ICHUZ.
A94067; ICGIZ.
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STRUCTURE BY NMR.
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S77834;
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PR00265; INTERLEUKIN2
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nes 30; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                        TISSUE-PETIPHERAL Blood;
Yabe M., Matsuura Y., Tatsumi M.;
Yabe M., Matsuura Y., Tatsumi M.;
Submitted (JUL-1995) to the EMBL/GenBank/DDBJ databases.

-!- FUNCTION: PRODUCED BY T-CELLS IN RESPONSE TO ANTIGENIC OR
MITOGRNIC STIMULATION, THIS PROTEIN IS REQUIRED FOR T-CELL
PROLIFERATION AND OTHER ACTIVITIES CRUCIAL TO REGULATION OF THE
IMMUNE RESPONSE. CAN STIMULARE B CELLS, MONCYTES, LYMPHOKINE-
ACTIVATED KILLER CELLS, NATURAL KILLER CELLS, AND GLIOMA CELLS (BY
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INTERCEDUKIN.2.
O-LINKED (GALNAC. .) (BY SIMILARITY).
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Mammalia; Butheria; Primates; Catarrhini; Cercopithecidae;
Cercopithecinae; Macaca.
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PROSITE; PS00424; INTERLEUKIN_2; 1.
Cytokine; Glycoprotein; Immune response; Signal; Growth factor;
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                                                                                                                                                                                                                                                                                       Macaca fascicularis (Crab eating macaque) (Cynomolgus monkey).
                                                                                                                                                                                                          01-NOV-1997 (Rel. 35, Created)
01-NOV-1997 (Rel. 35, Last sequence update)
16-CCT-2001 (Rel. 40, Last annotation update)
Interleukin-2 precursor (IL-2) (T-cell growth factor) (TGF).
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              100.0%; Score 147; DB 1; 100.0%; Pred. No. 2.5e-15;
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SUBCELLULAR LOCATION: Secreted.
SIMILARITY: BELONGS TO THE IL-2 FAMILY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ; Pred. No. 2.5
0; Mismatches
                                           0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1 APTSSSTKKTQLQLEHLLLDLQMILNGINN 30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   50
                                                                                         21 APTSSSTKKTOLOLEHLLLDLOMILNGINN 50
                                                                         1 APTSSSTKKTQLQLEHLLLDLQMILNGINN 30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      21 APTSSSTKKTQLQLEHLLLDLQMILNGINN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PRINTS; PR00265; INTERLEUKIN2.
ProDom; PD003649; Interleukin-2; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               InterPro; IPR000779; Interleukin-2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             154 AA; 17686 MW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             100.08;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              EMBL; D63352; BAA09676.1; -. HSSP; P01585; 3INK.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        100.08;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
                                           Conservative
                                                                                                                                                                               STANDARD;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   20
154
23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Pfam; PF00715; IL2; 1
                            Best_Local Similarity
Matches 30; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Local Similarity
                                                                                                                                                                                                                                                                                                                                                                SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                               NCBI_TaxID=9541;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       30;
                                                                                                                                                                              IL2_MACFA
Q29615;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DISULFID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CARBOHYD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
               Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       r-cell.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SIGNAL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Best Loca
Matches
                                                                                                                                                 RESULT 2
IL2_MACFA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                RESULT 3
IL2_MACMU
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             J. Immunol. 155:3946-3954(1995).

-i- FUNCTION: PRODUCED BY T-CELLS IN RESPONSE TO ANTIGENIC OR MITOGRAIC STIMULATION, THIS PROTEIN IS REQUIRED FOR T-CELL PROLIERATION AND OTHER ACTIVITIES CRUCIAL TO REGULATION OF THE IMMUNE RESPONSE. CAN STIMULATE B CELLS, MONOCYTES, LYMPHOKINE-ACTIVATED KILLER CELLS, NATURAL KILLER CELLS, AND GLIOMA CELLS (BY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               O-LINKED (GALNAC. .) (BY SIMILARITY).
BY SIMILARITY.
6AEBA480F204BA49 CRC64;
                                                                                                                                                                                      Macaca nemestrina (Pig-tailed macaque).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Cercopithecidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Villinger F.J., Brar S.S., Mayne A.E., Chikkala N., Ansari A.A.. "Comparative sequence analysis of cytokine genes from human and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SMART; SM00189; IL2; 1.
PROSITE; PS00424; INTERLEUKIN_2; 1.
Cytokine; Glycoprotein; Immune response; Signal; Growth factor;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ö
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01-NOV-1995 (Rel. 32, Last sequence update)
16-OCT-2001 (Rel. 40, Last annotation update)
Interleukin-2 precursor (IL-2) (T-cell growth factor) (TGF).
                           01-OCT-1996 (Rel. 34, Last sequence update)
16-OCT-2001 (Rel. 40, Last annotation update)
Interleukin-2 precursor (IL-2) (T-cell growth factor) (TCGF).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length 154;
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Pred. No. 2.5e-15;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               154 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      BY SIMILARITY. INTERLEUKIN-2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SIMILARITY).
-!- SUBCELLULAR LOCATION: Secreted.
-!- SIMILARITY: BELONGS TO THE IL-2 FAMILY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  21 APTSSSTKKTQLQLEHLLLDLQMILNGINN 50
                                                                                                                                                           Macaca mulatta (Rhesus macaque), and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         InterPro; IPR000779; Interleukin-2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PD003649; Interleukin-2; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                   MEDLINE-96003435; PubMed-7561102;
01-OCT-1996 (Rel. 34, Created)
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CARBOHYD
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                                                                                                                                                                                                                                                                                                                                                                                                                                                     Matches
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IL2_FELCA
   δ
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
                                                                                                                                          J. Immunol. 155:3946-3954(1995).
-1- FUNCTION: PRODUCED BY T-CELLS IN RESPONSE TO ANTIGENIC OR MITOGENIC STIMULATION, THIS PROTIEN SEQUIRED FOR T-CELL PROLIEDERATION AND OTHER ACTIVITIES CRUCIAL TO REGULATION OF THE IMMUNE RESPONSE. CAN STIMULATE B CELLS, MONOCYTES, LYMPHOKINE-ACTIVITIES CELLS, MONOCYTES, LYMPHOKINE-SCIPERELLUIAR LOCATION: Secreted.
-1- SUBCELLUIAR LOCATION: Secreted.
-1- SIMILARITY: BELONGS TO THE IL-2 FAMILY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SIMILARITY)
Cercocebus torquatus atys (Red-crowned mangabey) (Sooty mangabey).
Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi;
Mammalia, Eutheria, Primates, Catarrhini, Cercopithecidae,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Eukaryota; Metazoa; Chordata; Craniatà; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Carnivora; Pinnipedia; Phocidae; Mirounga.
                                                                                                              Ansari A.A.;
                                                                                                          Villinger F.J., Brar S.S., Mayne A.E., Chikkala N., Ansari A.A. "Comparative sequence analysis of cytokine genes from human and
                                                                                                                                                                                                                                                                                                                                                                                                                                    PROSITE; PS00424, INTERLEUKIN_2; 1.
Cytokine; Glycoprotein; Immune response; Signal; Growth factor;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ô
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             15-DEC-1998 (Rel. 37, Created)
15-DEC-1998 (Rel. 37, Last sequence update)
16-DCT-2001 (Rel. 40, Last annotation update)
Interleukin-2 precursor (IL-2) (T-cell growth factor) (TGF).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 154;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 142; DB 1; Length 15
Pred. No. 1.4e-14;
0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   .) (BY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               9FEB51814204BA48 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mirounga angustirostris (Northern elephant seal).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    INTERLEUKIN-2.
O-LINKED (GALNAC.
BY SIMILARITY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       154 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             BY SIMILARITY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE FROM N.A. MEDLINE=98136706; PubMed=9476229; Shoda L.K.M., Brown W.C., Rice-Ficht A.C.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1 APTSSSTKKTQLQLEHLLLDLQMILNGINN 30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  21 APTSRSTKKTQLQLEHLLLDLQMILNGINN 50
                                                                                                                                                                                                                                                                                                                                                                            InterPro; IPR000779; Interleukin-2. Pfam; PF00715; IL2; 1.
                                                                                                                                                                                                                                                                                                                                                                                                 PRINTS; PR00265; INTERLEUKINZ,
ProDom; PD003649; Interleukin-2; 1.
                                                                                              MEDLINE-96003435; PubMed-7561102;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             17754 MW;
                                                                                                                                                                                                                                                                                                                                                     EMBL; U19846; AAB60399.1; -.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     96.68;
96.78;
                                    Cercopithecinae; Cercocebus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       STANDARD;
                                                                                                                                                                                                                                                                                                                                                                                                                             SMART; SM00189; IL2; 1.
                                                                                                                                     nonhuman primates."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               154 AA;
                                                                                                                                                                                                                                                                                                                                                                  3 INK
                                                                      SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Similarity
                                               NCBI_TaxID-9531;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NCBI_TaxID=9716;
                                                                                                                                                                                                                                                                                                                                                                  HSSP; P01585;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              29;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      IL2_MIRAN
062641;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   VARIANT
SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CARBOHYD
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                                                                                                                                                                                                                                                                                                                                                                                                                                                               -cell,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SIGNAL
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                         J. Wildl. Dis. 34:81-90(1998).

-1- FUNCTION: PRODUCED BY T-CELLS IN RESPONSE TO ANTIGENIC OR MITOGENIC STHULLATION, THIS PROTEIN IS REQUIRED FOR T-CELL PROLIFERATION AND OTHER ACTIVITIES CRUCIAL TO REGULATION OF THE IMMUNE RESPONSE. CAN STIMULATE B CELLS, MONOCYTES, LYMPHOKINE-ACTIVATED KILLER CELLS, NATURAL KILLER CELLS, AND GLIOMA CELLS (BY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SIMILARITY).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         -i- FUNCTION: PRODUCED BY T-CELLS IN RESPONSE TO ANTIGENIC OR MITOGENIC STIMULATION, THIS PROTEIN IS REQUIRED FOR T-CELL PROLIFERATION AND OTHER ACTIVITIES CRUCIAL TO REGULATION OF THE IMMUNE RESPONSE. CAN STIMULATE B CELLS, MONCOYTES, LIMPHONINE- ACTIVATED KILLER CELLS, NATURAL KILLER CELLS, AND GLIOMA CELLS.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Felis silvestris catus (Cat).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Carnivora; Fissipedia; Felidae; Felis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Cytokine; Glycoprotein; Immune response; Signal; Growth factor; T-cell.
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01-0CT-1994 (Rel. 30, Last sequence update)
15-DEC-1998 (Rel. 37, Last annotation update)
Interleukin-2 precursor (IL-2) (T-cell growth factor) (TCGF).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 154;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE FROM N.A.
MEDLINE-93356765; PubMed-8352761;
Cozzi P.J., Padrid P.A., Takeda J., Alegre M.-A., Yuhki N.,
"Sequence and characterization of phocine interleukin 2.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 116; DB 1; Lengtu Lered. No. 1.3e-10;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE FROM N.A.
Litman R., Gibbs C., Good R.A., Day N.K.;
Submitted (NOV-1994) to the EMBL/GenBank/DDBJ databases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            . .) (BY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             INTERLEUKIN-2.
O-LINKED (GALNAC. . .) (B)
BY SIMILARITY.
OC92337A4B16B6BB CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            "Sequence and functional characterization of feline
Biochem. Biophys. Res. Commun. 194:1038-1043(1993).
                                                                                                                                                                                                                                           SUBCELLULAR LOCATION: Secreted. SIMILARITY: BELONGS TO THE IL-2 FAMILY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                154 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    BY SIMILARITY.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             78.9%; Score 116; 73.3%; Pred. No. 1
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| APTISSIKETQQQLEQLLLDLRLLLNGVNN 50
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PRINTS; PR00265; INTERLEUKIN2,
ProDom; PD003649; Interleukin-2; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      InterPro; IPR000779; Interleukin-2.
Pfam; PF00715; IL2; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PROSITE; PS00424; INTERLEUKIN_2; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        17661 MW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             EMBL; U79187; AAC12258.1; -.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                STANDARD;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 20
154
23
126
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                117;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        154 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ProDom; PD003649
SMART; SM00189;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NCBI_TaxID=9685
                                                                                                                                                                                                        SIMILARITY)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             HSSP; P01585;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         22;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Leff A.R.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             IL2_FELCA
Q07885;
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IL2_RABIT
                                                                                                                                                                                                                                                                                                                                                              CARBOHYD
CARBOHYD
DISULFID
                                                                                                                                                                                                                                                                                                                                                                                                                  CONFLICT
CONFLICT
SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
                                                                                                                                                                                                                                                                                                                                                                                                CONFLICT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        IL2_RABIT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              RESULT 8
 엄
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                            This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation-the Buropean Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/
                                                                                                                                                                                                                                                                                                                                                               ő
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               MEDLINE-96016696; PubMed-8571541;
Somberg R.L., Pullen R.P., Casal M.L., Patterson D.F., Felsburg P.J.,
Henthorn P.S.;
                                                                                                                                                                                                                                                                                                                                                                Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                "A single nucleotide insertion in the canine interleukin-2 receptor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Canis familiaris (Dog).
Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi,
Mammalia, Eutheria, Carnivora, Fissipedia, Canidae, Canis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             gamma chain results in X-linked severe combined immunodeficiency
                                                                                                                                                                                                                  PROSITE; PSO0424, INTERLEUKIN_2; 1.
Cytokine; Glycoprotein; Immune response; Signal; Growth factor;
                                                                                                                                                                                                                                                                                    .) (POTENTIAL)
                                                                                                                                                                                                                                                                                                                                                                ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                           1L2_CANFA STANDARD; PRT; 155 AA.
029416; 028249;
15-JUL-1998 (Rel. 36, Created)
15-JUL-1998 (Rel. 36, Last sequence update)
30-MAY-2000 (Rel. 39, Last annotation update)
Interleukin-2 precursor (IL-2) (T-cell growth factor) (TGGF).
                                                                                                                                                                                                                                                                                                                                          Length 154;
                                                                                                                                                                                                                                                                                                                                                               3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE FROM N.A.
STRAIN-XBRED21/12/93; TISSUE-Lymph node;
MEDLINE-95337423; Pubmed-7612930;
Dunham S.P., Argyle D.J., Onions D.E.;
"The isolation and sequence of canine interleukin-2.";
DNA Seq. 5:177-180(1995).
                                                                                                                                                                                                                                                                                                         -> I (IN REF. 2).
2E71E3BD8B9665EF CRC64;
                                                                                                                                                                                                                                                                                                                                         Score 114; DB 1;
Pred. No. 2.6e-10;
                                                                                                                                                                                                                                                                                    N-LINKED (GLCNAC.
KI -> RM (IN REF.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Vet. Immunol. Immunopathol. 47:203-213(1995)
                                                                                                                                                                                                                                                    BY SIMILARITY.
INTERLEUKIN-2.
BY SIMILARITY.
SUBCELLULAR LOCATION: Secreted. SIMILARITY: BELONGS TO THE IL-2 FAMILY.
                                                                                                                                                                                                                                                                                                                                                               5; Mismatches
                                                                                              or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                              1 APTSSSTKKTQLQLEHLLLDLQMILNGINN 30
                                                                                                                             EWEL; L25408; AAA51431.1; -.
PIR; JN0698; JN0698.
HSSP; P01585; 31NK.
InterPro; IPR000779; Interleukin-2.
Pfam; PR00715; IL2; 1.
PRINTS; PR00265; INTERLEUKIN2.
ProDom; PD003649; Interleukin-2; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE FROM N.A.
STRAIN-BEAGLE; TISSUE-Spleen;
MEDLINE-95347614; PubMed-7622066;
                                                                                                                                                                                                                                                                                                                   17653 MW;
                                                                                                                     EMBL; L19402; AAA02865.1; -.
                                                                                                                                                                                                                                                                                                                                         77.68;
73.38;
                                                                                                                                                                                                                                                                                                                                                               Conservative
                                                                                                                                                                                                          SMART; SM00189; IL2; 1.
                                                                                                                                                                                                                                                                                                         150 1
154 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            [2]
SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                    Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NCBI_TaxID=9615;
                                                                                                                                                                                                                                                              CHAIN
DISULFID
                                                                                                                                                                                                                                                                                             CONFLICT
CONFLICT
SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      disease."
                                                                                                                                                                                                                                                                                     CARBOHYD
                                                                                                                                                                                                                                                                                                                                          Query Match
                                                                                                                                                                                                                                        T-cell.
SIGNAL
                                                                                                                                                                                                                                                                                                                                                      Local
                                                                                                                                                                                                                                                                                                                                                               Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                  IL2_CANFA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  INTERLEUKIN-2.
O-LINKED (GALNAC. .) (BY SIMILARITY).
N-LINKED (GLCNAC. .) (POTENTIAL).
BY SIMILARITY.
M -> I (IN REF. 3).
O -> R (IN REF. 3).
F -> Y (IN REF. 3).
L -> M (IN REF. 3).
K -> Y (IN REF. 3).
K -> Y (IN REF. 3).
K -> M (IN REF. 3).
K -> M (IN REF. 3).
K -> M (IN REF. 3).
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                                                                                         -1- FUNCTION: PRODUCED BY T-CELLS IN RESPONSE TO ANTIGENIC OR MITOGENIC STIMULATION, THIS PROTEIN IS REQUIRED FOR T-CELL PROLIFERATION AND OTHER ACTIVITIES CRUCIAL TO REGULATION OF THE IMMUNE RESPONSE. CAN STIMULATE B CELLS, MONOCYTES, LYMPHOKINE-ACTIVATED KILLER CELLS, NATURAL KILLER CELLS, AND GLIOMA CELLS.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TISSUENCE FROM N.A.

TISSUE-Lymph node;

Kerr P.J., Lei S., Hardy C., Perkins H.D.;

Kerr P.J., Lei S., Hardy C., Perkins H.D.;

Kerr P.J., Lei S., Hardy C., Perkins H.D.;

Komplete cDNA sequence of rabbit interleukin-2.";

Submitted (MAY-1998) to the EMBL/GenBank/DDBJ databases.

-!- FUNCTION: PRODUCED BY T-CELLS IN RESPONSE TO ANTIGENIC OR

MITOGENIC STIMULATION, THIS PROTEIN IS REQUIRED FOR T-CELL

PROLIFERATION AND OTHER ACTIVITIES CRUCIAL TO REGULATION OF THE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Oryctolagus cuniculus (Rabbit).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Lagomorpha; Leporidae; Oryctolagus.
NCBI_TaxID=9986;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PROSITE; PS00424; INTERLEUKIN_2; 1.
Cytokine; Glycoprotein; Immune response; Signal; Growth factor;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ä
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length 155;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      15-JUL-1999 (Rel. 38, Created)
15-JUL-1999 (Rel. 38, Last sequence update)
16-OCT-2001 (Rel. 40, Last annotation update)
Interleukin-2 precursor (IL-2) (T-cell growth factor) (TCGF).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Indels
Knapp D.W., Williams J.S., Andrisani O.M.;
"Cloning of the canine interleukin-2-encoding cDNA.";
Gene 159:281-282(1995).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       73.1%; Score 107.5; DB 1
71.0%; Pred. No. 2.6e-09;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                153 AA
                                                                                                                                                                                                                                                                                       -!- SIMILARITY: BELONGS TO THE IL-2 FAMILY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          BY SIMILARITY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         6; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1 AP-TSSSTKKTQLQLEHLLLDLQMILNGINN 30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          21 APITSSSTKETEQOMEQLLLDLQLLLLNGVNN 51
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Pram; PR00715; IL2; 1.
PRINTS; PR00265; INTERLEUKIN2.
ProDom; PD003649; Interleukin-2; 1.
SMART; SM00189; IL2; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PRT;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              InterPro; IPR000779; Interleukin-2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              17668 MW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   EMBL; D30710; BAA06378.1; -.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       EMBL; U28141; AAA6969.1;
EMBL; U11689; AAA75360.1;
HSSP; P01585; 31NK.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STANDARD;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Cytokine; Glycoprotein;
T-cell.
SIGNAL 1 20
CHAIN 21 155
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       37
151
154
155 AA;
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Matches 22; Conserv
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entities requires a license agreement (See http://www.isb-sib.ch/announce/
                                     EMBL; M22899; AAA41427.1;
                                                                                                                                                                                                                                                                            Conservative
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                                                 PIR; A31278; A31278.
                                                                                                                                                                                                                                                                Similarity
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SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sus scrofa (Pig).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TISSUE-T-cell;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TISSUE-T-cell
                                                                                                                                                                                                                                                     Query ...
Best Local Similarian 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Lefevre F.
                                                                                                                                                                                                    CARBOHYD
DISULFID
SEQUENCE
                                                                                                                                                                                                                                                     Query Match
                                                                                                                                                                                                                                                                                                                                                                                    IL2_PIG
P26891;
                                                                                                                                                                   -cell.
                                                                                                                                                                               SIGNAL
                                                                                                                                                                                                                                                                                                                                                              RESULT 10
  6
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                                                                                                                                                                                                                                                                                                                                                                                                  This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics. Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation-the Buropean Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial
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                                                                                                                                                                                                                                                                                                                                                                                     Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Rattus.
                                                                                                                                                                                                                                                                                                O-LINKED (GALNAC. . .) (BY SIMILARITY).
N-LINKED (GLCNAC. . .) (POTENTIAL).
BY SIMILARITY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             IMMUNE RESPONSE. CAN STIMULATE B CELLS, MONOCYTES, LYMPHOKINE-ACTIVATED KILLER CELLS, AND GLIOMA CELLS.
-1- SUBCELLULAR LOCATION: Secreted.
-1- SIMILARITY: BELONGS TO THE IL-2 FAMILY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                MEDLINE=89339608; PubMed=2788130;
MCKnight A.J., Mason D.W., Barclay A.N.;
"Sequence of rat interleukin 2 and anomalous binding of a mouse
interleukin 2 cDNA probe to rat MHC class II-associated invariant
                                                                                                                                                                                                                                      PROSITE; PS00424; INTERLEUKIN_2; 1.
Cytokine; Glycoprotein; Immune response; Signal; Growth factor;
                                                                                                                                                                                                                                                                                                                                                                                    ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                01-AUG-1990 (Rel. 15, Created)
01-AUG-1990 (Rel. 15, Last sequence update)
16-OCT-2001 (Rel. 40, Last annotation update)
Interleukin-2 precursor (IL-2) (T-cell growth factor) (TCGF).
                                                                                                                                                                                                                                                                                                                                                            Length 153;
                                                                                                                                                                                                                                                                                                                                                Score 107; DB 1; Length 13.
Pred. No. 3.1e-09;
                                                                                                                                                                                                                                                                                                                          SIMILARITY.
8173536B2DDD8B86 CRC64;
                                                                                                                                                                                                                                                                          BY SIMILARITY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             155 AA
                                                                                                                                                                                                                                                                                      INTERLEUKIN-2.
                                                                                                                                                                                                                                                                                                                                                                                                                       21 APTSSSTKETQEQLDQLLLDLQVLLKGVND 50
                                                                                                                                                                                                                                                                                                                                                                                                          1 APTSSSTKKTQLQLEHLLLDLQMILNGINN 30
                                                                                                                                                                  HSSP; P01585; 31NK.
InterPro; IPR000779; Interleukin-2.
Pfam; PF00715; IL2; 1.
                                                                                                                                                                                                    PRINTS; PR00265; INTERLEUKIN2.
ProDom; PD003649; Interleukin-2; 1.
SMART; SM00189; IL2; 1.
                                                                                                                                                                                                                                                                                                                                                                                   ;9
                                                                                                                                                      EMBL; AF068057; AAC23838.1; -.
                                                                                                                                                                                                                                                                                                                                     17256 MW;
                                                                                                                                                                                                                                                                                                                                                            72.8%;
70.0%;
                                                                                                                                                                                                                                                                                                                                                           Query Match 72.8%
Best Local Similarity 70.0%
Matches 21; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                IL2 OR IL-2.
Rattus norvegicus (Rat).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            STANDARD;
                                                                                                                                                                                                                                                                                                                                     153 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NCBI_TaxID=10116;
                                                                                                                                                                                                                                                                                                                         DISULFID
SEQUENCE
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                                                                                                                                                                                                                                                                                                             CARBOHYD
                                                                                                                                                                                                                                                              T-cell.
SIGNAL
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Goodall J.C., Emery D.C., Bailey M., English L.S., Hall L.; "CDNA cloning of porcine interleukin 2 by polymerase chain reaction."; Biochim. Biophys. Acta 1089:257-258(1991).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                             O-LINKED (GALNAC. . .) (BY SIMILARITY).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Cetartiodactyla; Suina; Suidae; Sus.
NCBI_TaxID=9823;
                                                                                                                                                                                                                                                                                          PROSITE; PS00424; INTERLEUKIN_2; 1.
Cytokine; Glycoprotein; Immune response; Signal; Growth factor;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       01-AUG-1992 (Rel. 23, Created)
01-AUG-1992 (Rel. 23, Last sequence update)
01-MAR-2002 (Rel. 41, Last annotation update)
Interleukin-2 precursor (IL-2) (T-cell growth factor) (TCGF).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     66.0%; Score 97; DB 1; Length 155; 66.7%; Pred. No. 1e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          6; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Submitted (MAY-1991) to the EMBL/GenBank/DDBJ databases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  67A8554A73BF30A0 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          154 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          4; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                    INTERLEUKIN-2
                                                                                                                                                                                                                                                                                                                                                                                                                                                            BY SIMILARITY
or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1 APTSSSTKKTQLQLEHLLLDLQMILNGINN 30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PIR; A45882; A45882.
HSSP, P01585; 31NK.
InterPro; IPR000779; Interleukin-2.
Pfam; PF00715; IL2; 1.
PRIMTS; PR00265; INTERLEUKIN2.
ProDom; PD003649; Interleukin-2; 1.
SMART; SM00189; IL2; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   MEDLINE~91274360; PubMed=2054386;
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IPR000779; Interleukin-2.

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entities requires a license agreement (See http://www.isb-sib.ch/announce/
or send an email to license@isb-sib.ch).
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"Cross-species PCR cloning of garbil (Mariones unquiculatus)
interleukin-2 cDNA and its expression in COS-7 cells.";
Vet. Immunol. Immunopathol. 40:63-71(1994).
-!- FUNCTION: PRODUCED BY T-CELLS IN RESPONSE TO ANTIGENIC OR
MITOGENIC STIMULATION, THIS PROTIEN SEQUIRED FOR T-CELL
PROLIFERATION AND OTHER ACTIVITIES CRUCIAL TO REGULATION OF THE
IMMUNE RESPONSE. CAN STIMULATE B CELLS, MONOCYTES, LYMPHOKINE-
ACTIVATED KILLER CELLS, NATURAL KILLER CELLS, AND GLIOMA CELLS.
-!- SUBCELLULAR LOCATION: Secreted.
-!- SIMILARITY: BELONGS TO THE IL-2 FAMILY.
                                                                                                                                                                                                                                                                                                                        . .) (BY SIMILARITY).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Gerbillinae;
                                                                                                                                                                                                                                                   Cytokine; Glycoprotein; Immune response; Signal; Growth factor;
                                                                                                                                                                                                                                                                                                                                                                                                                          ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         01-OCT-1994 (Rel. 30, Last sequence update)
16-OCT-2001 (Rel. 40, Last annotation update)
Interleukin-2 precursor (IL-2) (T-cell growth factor) (TCGF).
                                                                                                                                                                                                                                                                                                                                                                                          Length 154;
                                                                                                                                                                                                                                                                                                                                                                         Score 96; DB 1; Length 134
Pred. No. 1.5e-07;
Trefile 154
                                                                                                                                                                                                                                                                                                                                                      F3B95E43D4A3D3E1 CRC64;
                                                                                                                                                                                                                                                                                                                    O-LINKED (GALNAC.
BY SIMILARITY.
                                                                                                                                                                                                                                                                                      BY SIMILARITY
                                                                                                                                                                                                                                                                                                                                                                                                                        4; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Meriones unguiculatus (Mongolian jird).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         21 APTSSSTKNTKKQLEPLLLDLQLLLKEVKN 50
                                                                                                                                                                                                                                                                                                                                                                                                                                                        1 APTSSSTKKTQLQLEHLLLDLQMILNGINN 30
                                                                             EMBL; AB041935; BAB16110.1; -. PIR; S15473; S15473. BAB16110.1; -. PIR; S16241; S16241. HASSP; P01565; 31NK. InterPro; IPR000779; Interleukin-2.
                                                                                                                                                             Pfam; PF00715; IL2; 1.
PRINTS; PR00265; INTERLEUKIN2.
ProDom; PD003649; Interleukin-2; 1.
SMART; SM00189; IL2; 1.
                                                                                                                                                                                                                  SMART; SM00189; IL2; 1.
PROSITE; PS00424; INTERLEUKIN_2; 1.
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MEDLINE-94174702; PubMed-8128610;
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(Rel. 30, Last sequ
(Rel. 40, Last anno
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154 IN
23 O-
126 BY
17401 MW;
                                               EMBL; X56750; CAA40071.1; -.
                                                                                                                                                                                                                                                                                                                                                                                     65.3%;
ilarity 66.7%;
Conservative
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                                                                  EMBL; X58428; CAA41330.1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            STANDARD;
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                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local Similarity
Matches 20; Conserv
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78
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01-0CT-1994
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Q08081;
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                                                                                                                                                                                                                                                                                                                                                       SEQUENCE
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                                                                                                                                                                                                                                                                      Gaps
                                                                                                                                                                   . .) (BY SIMILARITY).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PROLIEBRATION AND OTHER ACTIVITIES CRUCIAL TO REGULATION OF THE IMMUNE RESPONSE. CAN STIMULATE B CELLS, MONOCYTES, LYMPHOKINE-ACTIVATED KILLER CELLS, AND GLIOMA CELLS. SUBCELLULAR LOCATION: Secreted. SIMILARITY: BELONGS TO THE IL-2 FAMILY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi,
Mammalia, Eutheria, Perissodactyla, Equidae, Equus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PRINTS; PR00265; INTERLEUKIN2.
Prodom; PD003649; Interleukin-2; 1.
SMART; SM0189; IL2; 1.
PROSTIE; PS00424; INTERLEUKIN.2; 1.
CYtokine; Glycoprotein; Immune response; Signal; Growth factor;
                                                            PROSITE; PS00424; INTERLEUKIN_2; 1.
Cytokine; Glycoprotein; Immune response; Signal; Growth factor; T-cell.
SIGNAL.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Tavernor A.S., Allen W.R., Butcher G.W.;
Submitted (NOV-1992) to the EMBL/GenBank/DDBJ databases.
-!- FUNCTION: PRODUCED BY T-CELLS IN RESPONSE TO ANTIGENIC OR
MITOGENIC STIMULATION, THIS PROTEIN IS REQUIRED FOR T-CELL
                                                                                                                                                                                                                                                                                                                                                                                                                                                       01-0cT-1994 (Rel. 30, Created)
01-NOV-1995 (Rel. 32, Last sequence update)
16-0cT-2001 (Rel. 40, Last annotation update)
Interleukin-2 precursor (IL-2) (T-cell growth factor) (TGF).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                MEDLINE-94160538; PubMed-8116217; Vandergrifft E.V., Horohov D.W.; Molecular cloning and expression of equine interleukin 2."; Vet. Immunol, Immunopathol. 39:395-406(1993).
                                                                                                                                                                                                                                  Score 95; DB 1; Length 155;
Pred. No. 2.1e-07;
2; Mismatches 8; Indels
                                                                                                                                                   INTERLEUKIN-2.
O-LINKED (GALNAC. . .) (BY
BY SIMILARITY.
DOF74AA1A381CDDA CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                           149 AA.
                                                                                                                                                                                                                                                                    2; Mismatches
                                                                                                                                                                                                                                                                                                      1 APTSSSTKKTQLQLEHLLLDLQMILNGINN 30
                                                                                                                                                                                                                                                                                                                         21 APTSSPAKEAQOYLEQLLLDLQQLLRGINN 50
            Pfam; PF00715; IL2; 1.
PRINTS; PR00765; INTERLEUKIN2.
PP0DOM; PD003649; INTErleukin-2; 1.
SMART; SM00189; ILZ; 1.
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Pfam; PF00715; IL2; 1.
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155 11
23 O
126 B1
17602 MW;
                                                                                                                                                                                                                                  64.68;
66.78;
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                                                                                                                                                                                                                                                                    Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                           STANDARD;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Equus caballus (Horse)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PIR; S31391; S31391.
HSSP; P01585; 3INK.
                                                                                                                                               21 1
23 78 1
155 AA;
                                                                                                                                                                                                                                                      Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE FROM N.A.
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                                                                                                                                                                   CARBOHYD
                                                                                                                                                                                   DISULFID
                                                                                                                                                                                                    SEQUENCE
                                                                                                                                                                                                                                      Query Match
                                                                                                                                                                                                                                                      Local
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                                                                                                                                                    CHAIN
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CHAIN
CARBOHYD
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NON_TER
SEQUENCE
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-1- FUNCTION: PRODUCED BY T-CELLS IN RESPONSE TO ANTICENIC OR MITOGENIC STIMULATION, THIS PROTEIN IS REQUIRED FOR T-CELL PROLIFERATION AND OTHER ACTIVITIES CRUCIAL TO REGULATION OF THE IMMUNE RESPONSE. CAN STIMULATE B CELLS, MONOCYTES, LYMPHOKINE-ACTIVATED KILLER CELLS, NATURAL KILLER CELLS, AND GLIOMA CELLS (BY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Orcinus orca (Killer whale).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Cetartiodactyla; Cetacea; Odontoceti; Delphinidae;
                                                                                                                                                                                                                                                                                                                                 Gaps
                                                                          . .) (BY SIMILARITY).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Ness T.L., Bradley W.G., Reynolds J.E. III, Roess W.B.; "Isolation and expression of the interleukin-2 gene from the killer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PROSITE; PS00424; INTERLEUKIN 2; 1.
Cytokine; Glycoprotein; Immune response; Signal; Growth factor; T-cell.
SIGNAL 1 20 RY SIMITADITOV
                                                                                                                                                                                                                                                                                                                               ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        30-WAY-2000 (Rel. 39, Created)
30-WAY-2000 (Rel. 39, Last sequence update)
16-CCT-2001 (Rel. 40, Last annotation update)
Interleukin-2 precursor (IL-2) (T-cell growth factor) (TCGF)
                                                                                                                                                                                                                                                                                           Length 149;
                                                                                                                                                                                                                                                        051BB8C47A0114FC CRC64;
                                                                                                                                                                                                                                                                                         DB 1,
5.7e-07;
5;
                   BY SIMILARITY,
INTERLEUKIN-2.
BY SIMILARITY.
O-LINKED (GALNAC.
N-LINKED (GLCNAC.
R -> K (IN REF. 2).
I -> M (IN REF. 2).
I -> M (IN REF. 2).
E -> G (IN REF. 2).
I -> F (IN REF. 2).
I -> M (IN REF. 2).
I -> K (IN REF. 2).
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REF. 2).
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                                                                                                                                                                                                                                                                                                                                   8; Mismatches
                                                                                                                                                                                                                                                                                           62.6%; Score 92; 56.7%; Pred. No.
                                                                                                                                                                                                                                                                                                                                                                      1 APTSSSTKKTQLQLEHLLLDLQMILNGINN 30
                                                                                                                                                                                                                                                                                                                                                                                        PRODUCTOR TO THE STREET OF THE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PRT;
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HSSP; P01585; 1IRL.
                                                                                                                                                                                                                                                        17086 MW;
                                                                                                                                                                                                                                                                                                             Best Local Similarity 56.7
Matches 17; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        STANDARD;
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                       20
149
121
23
106
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8
125
125
145
148
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097513;
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CONFLICT
CONFLICT
CONFLICT
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                                                             DISULFID
                                                                                                                                                                                                                                                        SEQUENCE
                                                                                                                                                                                                                                                                                               Query Match
                                                                                CARBOHYD
                                                                                                      CARBOHYD
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       T-cell.
                                             CHAIN
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IL2_ORCOR
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INTERLEUKIN-2.
O-LINKED (GALMAC. . .) (BY SIMILARITY).
BY SIMILARITY.
                                                                                                                                                                                                                                                                 Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Anikeeva N.V., Vinogradova T.V., Votoshin O.N.;
Submitted (DEC-1989) to the EMBL/GenBank/DDBJ databases.
-! FUNCTION: PRODUCED BY T-CELLS IN RESPONSE TO ANTIGENIC OR
MITGERIC STIMULATION. THIS PROTEIN IS REQUIRED FOR T-CELL
PROLIFERATION AND OTHER ACTIVITIES CRUCIAL TO REGULATION OF THE
IMMUNE RESPONSE. CAN STIMULATE B CELLS, MONOCYTES, LYMPHOKINE-
ACTIVATED KILLER CELLS, NATURAL KILLER CELLS, AND GLIOMA CELLS.
-! SUBCELLULAR LOCATION: Secreted.
-! SIMILARITY: BELONGS TO THE IL-2 FAMILY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Bos taurus (Bovine).
Eukaryota, Metazoa, Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Cetartiodactyla; Ruminantia; Pecora; Bovoidea;
Bovidae; Bovinae; Bos.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE FROM N.A.
MEDLINE=86205870; PubMed=3486415;
Reeves R., Spies A.G., Nissen M.S., Buck C.D., Weinberg A.D.,
Barr P.J., Magnuson N.S., Magnuson J.A.;
"Molecular cloning of a functional bovine interleukin 2 cDNA.";
Proc. Natl. Acad. Sci. U.S.A. 83:3228-3232(1986).
                                                                                                                                                                                                                                                             ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ILZ_BOVIN STANDARD; PRT; 155 AA.
p05016.
13-AUG-1987 (Rel. 05, Created)
13-AUG-1987 (Rel. 05, Last sequence update)
16-0CT-2001 (Rel. 40, Last annotation update)
1nterleukin-2 precursor (IL-2) (T-cell growth factor) (TGF).
ILZ OR ILJ-2
                                                                                                                                                                                                    Length 152;
                                                                                                                                                                                                                                                                 Indels
               >152 INTERLEUKIN-2.
23 O-LINKED (GALNAC. . .) (BY
126 BY SIMILARITY.
152 IN 308F91821ECCB764 CRC64;
                                                                                                                                                                                                                                                             9
                                                                                                                                                                                              Score 88; DB 1; 1
Pred. No. 2.4e-06;
                                                                                                                                                                                                                                                                 Mismatches
                                                                                                                                                                                                                                                                                                                                                                   21 APTSSSTENTKKQVQSLLQDLQLLKEINN 50
                                                                                                                                                                                                                                                                                                                             1 APTSSSTKKTQLQLEHLLLDLQMILNGINN 30
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   EMBL; M12791; AAA30586.1; --
EMBL; M13204; AAA21143.1; ALT_INIT.
EMBL; X17201; CAA35062.1; --
EMBL; X22687; CAA36912.1; --
HSSP; P01585; 3INK.
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ProDom; PD003649; Interleukin-2; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MEDLINE-86205869; PubMed-3517854;
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                                                                                                                                                                                                                                                                    Conservative
                   21 >1
23
78 1
152 1
                                                                                                                                                                                                                              Local Similarity
nes 18; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NCBI_TaxID=9913;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              FISSUE-Thymus;
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                                                                                                                                                                                                                                                                                                                                            Gaps
                                                                                                              INTERLEUKIN-2.
BY SIMILARITY.
O-LINKED (GALNAC. . .) (BY SIMILARITY).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Cetartiodactyla; Ruminantia; Pecora; Bovoidea;
                                                   Cytokine; Glycoprotein; Immune response; Signal; Growth factor; T-cell.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PROSITE; PS00424; INTERLEUKIN_2; 1.
Cytokine; Glycoprotein; Immune response; Signal; Growth factor;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            112_CAPH1 STANDARD; PRT; 155 AA.
P36835; P79156;
01-JUN-1994 (Rel. 29, Last sequence update)
01-JUN-1994 (Rel. 29, Last sequence update)
16-OCT-2001 (Rel. 40, Last annotation update)
Interleukin-2 precursor (IL-2) (T-cell growth factor) (TCGF).
                                                                                                                                                                                                                                                                                      Length 155;
                                                                                                                                                                                                                                                                              / Match 53.7%; Score 79; DB 1; Length 155 Local Similarity 53.3%; Pred. No. 5.7e-05; les 16; Conservative 6; Mismatches 8; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Rimstad E.;
Submitted (NOV-1993) to the EMBL/GenBank/DDBJ databases.
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                                                                                                                                                                                                                                                                                                                                                                                                                21 APTSSSTGNTMKEVKSLLLDLQLLLERVKN 50
                                                                                                                                                                                                                                                                                                                                                                                          1 APTSSSTKKTQLQLEHLLLDLQMILNGINN 30
                        PROSITE; PS00424; INTERLEUKIN_2; 1.
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EMBL; U34274; AAB38527.1; -.
PIR; S38662, S38662.
HSSP, 01586; JINK.
InterPro; IPR000779; Interleukin-2.
Pfam; PF00715; IL2; 1.
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ProDom; PD003649; Interleukin-2; 1.
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155 IN
127 BX
23 O-
66 V
17627 MW;
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NCBI_TaxID=9925;
SMART; SM00189; IL2; 1
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155
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TISSUE-Blood;
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Gaps
    . .) (BY SIMILARITY).
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                                                                              (IN REF. 2).
                                                                                                                               Length 155;
                                                                                                                              Score 79; DB 1; Length 155
Pred. No. 5.7e-05;
6; Mismatches 8; Indels
                                                                                    0 -> L (IN REF. 2).
T -> I (IN REF. 2).
L -> M (IN REF. 2).
90022DFBB6AF78DE CRC64;
                     5
  O-LINKED (GALNAC. . BY SIMILARITY.

BY SIMILARITY.

P -> T (IN REF. 2).

T -> P (IN REF. 2).

T -> P (IN REF. 2).

D -> A (IN REF. 2).

MASLKG -> SMDNIKR (.)
53.7%;
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    CARBOHYD
           DISULFID
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    STREETHERE
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Search completed: October 21, 2002, 09:49:03 Job time : 12 secs

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GenCore version 5.1.3
Copyright (c) 1993 - 2002 Compugen Ltd.
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OM protein - protein search, using sw model

October 21, 2002, 09:47:56; Search time 16 Seconds (without alignments) 180.168 Million cell updates/sec Run on:

US-09-720-828A-4

1 APTSSSTKKTQLQLEHLLLDLQMILNGINN 30 Perfect score: Sequence:

BLOSUM62 Gapop 10.0 , Gapext 0.5 Scoring table:

283138 seqs, 96089334 residues Searched: Total number of hits satisfying chosen parameters:

Minimum DB seq length: 0 Maximum DB seq length: 200000000

Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 su

PIR_71:*
1: pir1:*
2: pir2:*
3: pir3:*
4: pir4:* Database

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

	Description	interleukin-2 pres	ď	interleukin 2 prec	interleukin-2 prec	interleukin-2 prec	interleukin-2 - Mo	interleukin-2 prec	interleukin-2 go	interleukin-2 prec	interleukin-2 prec	interleukin-2 prec	interleukin-2 prec	~		interleukin 2 - mo	isopenicillin N ac	probable exopolysa	probable LysR-type	flagellar hook-ass	hypothetical prote	methyl-accepting c	protein F52E1.4 [i	hypothetical prote				aminopeptidase C [pol polyprotein -	hypothetical prote
SUMMARIES	Œ	ICHU2	ICG12	JN0698	A31278	S16241	833509	S31391	S38662	S11488	145913	837289	ICMS2	168870	154512	168871	S12169	G82262	F95285	E70122	C70701	F82217	A89130	T22210	F64145	H64307	AB1367	AC1736	GNL,JG4	T11685
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_	Query Match	100.0	100.0	77.6	0.99	55.3	64.6	52.6	53.7	53.7	33.7	6.9	13.5	40.5	8.6	39.1	26.7	35.4	34.7	34.7	34.0	33.3	33.3	33.0	32.7	32.7	32.7	32.7	7.7	32.3
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polymorphic outer	polymorphic membra	DNA-binding respon	conserved hypothet	hypothetical prote	SEC14 protein homo	nonmuscle myosin I	hypothetical prote	hypothetical prote	hypothetical prote	hypothetical prote	galactokinase (EC	oligopeptide ABC t	peptide ABC transp	hypothetical prote	hypothetical prote
F86548	H72074	H82447	B70209	F89839	T41244	A59282	T05656	C84888	A71946	H90434	C37760	B64551	H82355	T18745	T32980
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82	33	32	32	32	E	32	3	31	3	33	31	33	33	33	33
47.5	47.5	47	47	47	47	47	46.5	46.5	46	46	46	46	46	46	46
30	31	32	33	34	35	36	37	38	39	40	41	42	43	44	45

ALIGNMENTS

RESULT 1

ICHU2
interleukin-2 precursor (validated) - human
interleukin-2 precursor [validated] - human N,Alternate names: IL-2; T-cell growth factor
interleukin-2 precursor (validated) - human N.Alternate names: IL-2; T-cell growth factor C.Species: Homo sapiens (man)
<pre>interleukin-2 precursor (validated] - human N.Alternate names: II2; T-cell growth factor C;Species: Homo sapiens (man) C;Species: 11-Aug-1983 #sequence_revision 11-Aug-1983 #text_cha</pre>
<pre>interleukin-2 precursor (validated] - human N.Alternate names: IL-2; T-cell growth factor C;Species: Homo sapiens (man) C;Species: Homo sapiens (man) C;Date: Il-Aug-1983 #sequence_revision 11-Aug-1983 #text_cha C;Date: Il-Aug-1983 #sequence_revision 11-Aug-1983 #text_cha C;Accession: A01849; A21192; A20961; S31209; A93297; A90113;</pre>

ange 08-Dec-2000 ; A93478; I56518; I73624

Ridibrook, N.J.; Lieber, M.; Crabtree, G.R.
Nucleic Acids Res. 12, 5005-5013, 1984
A.Title: DNA sequence of the 5' flanking region of the human interleukin 2 gene: ho
A; Reference number: A93524; MUID:84247353
A; Reference number: A01849
A; Molecule type: DNA
A; Molecule type: DNA
A; Residues: 1-153 <HOL>
A; Residues: 1-153 <HOL>
A; Cross-references: GB:X00695; GB:X00200; GB:X00201; GB:X00202; NID:933783; PIDN:CA
A; Cross-references: GB:X00695; GB:X00200; GB:X00201; GB:X00202; NID:933783; PIDN:CA
A; Cross-reference number: A0192; MUID:84170243
A; Title: Structure of the human interleukin 2 gene.

A; Accession: Manager A 21192
A; Molecule type: DNA
A; Molecule type: DNA
A; Residues: 1-153 < FUJ>
A; Cossion: 1-153 < FUJ>
A; Cossion: 1-153 < FUJ>
B; Goldon, I, J; Smith, K.A.; Fornace Jr., A.J.; Comeau, C.M.; Wiskocil, R.L.; Crah
Proc. Natl. Acad. Sci. U.S.A. 81, 1634-1638, 1984
A; Title: T-cell growth factor: complete nucleotide sequence and organization of the A; Reference number: A20961; MUID:84170356

A Molecule type: DNA
A; Molecule type: 1-153 -400>
A; Residues: 1-153 -400>
A; Cross-references: GB: KR02056; NID:9186302; PIDN: AAA98792.1; PID:9386819
B; Laabi, Y.; Gras, M.P.; Carbonnel, F.; Brouet, J.C.; Berger, R.; Larsen, C.J.; Tsag EMBO J. 11, 3897-3904, 1992
B; Labbo J. 11, 3897-3904, 1992
A; Title: A 'new gene, BCM, on chromosome 16 is fused to the interleukin 2 gene by a t A; Reference number: S31208; MUID:93010984

A; Accession: S31209
A; Molecule type: mRNA
A; Residues: 11-17 (LAA>
A; Cossiones: I1-117 (LAA>
A; Cossiones: EMBL:214955
A; Cross-references: EMBL:214955
A; Mote: this sequence is shown from the beginning of the fragment to the chromosomal A; Mote: this sequence is shown from the beginning of the fragment to the chromosomal B; Taniguchi, T.; Matsui, H.; Fujita, T.; Takaoka, C.; Kashima, N.; Yoshimoto, R.; Hai Nature 302, 305-310, 1983
A; Title: Structure and expression of a cloned cDNA for human interleukin-2.
A; Reference number: A93297; MUID:83167472

A; Molecule type: mRNA A;Residues: 1.153 TRAN> A;Cross-treferences: GR:V00564; MID:933780; PIDN:CAA23827.1; PID:933781 A;Experimental source: leukemic T-cell line, Jurkat-111, cloned from Jurkat-FHCRC R;Maded, S.; Nishino, N.; Obaru, K.; Mita, S.; Nomiyama, H.; Shimada, K.; Fujimoto, Biochem. Biophys: Res. Commun. 115, 1040-1047, 1983 A;Title: Cloning of interleukin 2 mRNAs from human tonsils.

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A; Accession: A34463
A; Molecule type: protein
A; Residues: 21-35 <CON>
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A; Status: translated from GB/EMBL/DDBJ
A; Status: translated from GB/EMBL/DDBJ
A; Molecule type: DNA
A; Residues: 1-68 - RE2>
A; Cross-references: GB:M31199; NID:g186296; PIDN:AAA59139.1; PID:g553508
A; Cross-references: GB:M31199; NID:g186296; PIDN:AAA59139.1; PID:g553508
B; Siebenlist, U.; Durand, D.B.; Bressler, P.; Holbrook, N.J.; Norris, C.A.; Kamoun, M.; Mol. Cell. Blol. 6, 3042-3049, 1986
A; Title: Promoter region of interleukin-2 gene undergoes chromatin structure changes and A; Reference number: I57603; MUID:87064618
A; Reference mumber: I57603
A; Status: translated from GB/EMBL/DDBJ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       9
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A; Note: heterogeneity in Jurkat-derived IL-2 is primarily due to differences in glycosyl
n in lacking 21-Ala (FT-112-A and FT-1L2-B) and 22-Pro (FT-IL2-B)
R; Conradt, H.S.; Nimtz, M.; Dittmar, K.E.J.; Lindenmaier, W.; Hoppe, J.; Hauser, H.
J. Biol. Chem. 264, 17368-17373, 1989
A; Title: Expression of human interleukin-2 in recombinant baby hamster kidney, Ltk-, and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       A; Status: translated from GB/EMBL/DDBJ
A; Status: translated from GB/EMBL/DDBJ
A; Status: translated from GB/EMBL/DDBJ
A; Status: translated by the state of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      expression in Escherichia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 A.Molecule type: mRNA
A.Residues: 1-153 <DEV>
A.A.Cross-references: GB-V00564; NID:g33780; PIDN:CAA23827.1; PID:g33781
A.Experimental source: splencytes
R.Eizenberg, O.; Faber Elman, A.; Lotan, M.; Schwartz, M.
A. Neurochem. 64, 1928-1936, 1995
A.7 Title: Interleukin-2 transcripts in human and rodent brains: possible expression by A.Reference number: 156518; MUID:95239150
A.Accession: 156518
A.Accession: 156518
A.Status: translated from GB/EMBL/DDBJ
A.Molecule type: mRNA
A.Residues: 1-152 <EIZ.
                                                                                                                                                                                                                                                                                                                                                   Degrave, W.; Tavernier, J.;
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A; Residues: 1-68 (RE3>
A; Cross-references: GB:M13879; NID:g186305; PIDN:AAA59141.1; PID:g553509
A; Cross-references: GB:M13879; NID:g186305; PIDN:AAA59141.1; PID:g553509
B; Weir, M.P.; Chaplin, M.A.; Wallace, D.M.; Dykes, C.W.; Hobden, A.N.
Biochemistry 27, 6883-6892, 1988
A; Title: Structure-activity relationships of recombinant human interleukin
A; Reference number: IS2401; Mull:89064420
A; Contents: recombinant IL-2 and mutants expressed in E. coli
                                                                                                             A; Molecule type: mRNA
A; Residues: 1-153 < MAE>
A; Cross-references: GB:J00264; NID:g186294; PIDN:AAD48509.1; PID:g5729676
A; Experimental source: consillar mononuclear cells
R; Devos, R.; Plaetinck, G.; Cheroutre, H.; Simons, G.; Degrave, W.; Tavern
Nucleic Acids Res. 11, 4307-4323, 1983
A; Title: Molecular cloning of human interleukin 2 cDNA and its expression
A; Reference number: A93478; MUID:83246551
A; Reference number: A90113; MUID: 84023840
A; Accession: A90113
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A; Residues: 21-153 <ROB>
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A) Note: the O-linked glycosylation site in recombinant material matched that from hur R; Grabenhorst, E.; Hofer, B.; Nimtz, M.; Jaeger, V.; Conradt, H.S.
Eur. J. Biochem. 215, 189-197, 1993
A; Title: Biosynthesis and secretion of human interleukin 2 glycoproteins variants fr A; Reference number: S34052; MUID:93345493
A; Contents: annotation; glycosylation of variant forms expressed in insect cells
                                                                                                                                                                                                                                                                                                                                                                                                                                                      A map position: 4q26-4q27

A; Interns: 49/3; 69/3; 117/3

C; Superfamily: interleukin-2
F; 1-20/Domain: signal sequence *status predicted <SIG>
F; 1-30/Domain: signal sequence *status experimental <IL2>
F; 23/Binding site: carbohydrate (Phr) (covalent) *status experimental
F; 78-125/Disulfide bonds: *status experimental
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       interleukin-2 precursor - common gibbon
NiAlternate names: IL-2; r-cell growth factor
C; Species: Hylobates lar (common gibbon, white-handed gibbon)
C; Date: 31-Dec-1991 #sequence_revision 31-Dec-1991 #text_change 22-Jun-1999
C; Accession: A94067; A01849
R; Chen, S.J.; Holbrook, N.J.; Mitchell, K.F.; Vallone, C.A.; Greengard, J.S.; Crabtr Proc. Natl. Acad. Sci. U.S.A. 82, 7284-7288, 1985
A; Title. A viral long terminal repeat in the interleukin 2 gene of a cell line that A; Reference number: A94067; MUID:86042650
A; Molecule type: mRNA
A; Residues: 1-153 CHE>
A; Molecule type: mRNA
A; Residues: 1-153 CHE>
A; Cross-references: GB:MIL144; NID:g177014; PIDN:AAA35454.1; PID:g177015
A; Experimental source: leukemia cell line MLA 144; ATCC TIB 201
A; Note: the integration of a retrovirus sequence containing a 5' LTR into the 3' non C; Superfamily: interleukin-2
C; Superfamily: interleukin-2
C; Keywords: cytokine; glycoprotein; growth factor; immunoregulation; lymphokine; T-C; Keywords: cytokine; glycoprotein; growth factor; immunoregulation; lymphokine; T-C; Keywords: cytokine; glycoprotein; growth factor; fumunoregulation; lymphokine; T-C; Keywords: cytokine; glycoprotein; growth factor; fumu
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R;Cozzi, P.J.: Padrid, P.A.; Takeda, J.; Alegre, M.L.; Yuhki, N.; Leff, A.R. Biochem. Biophys. Res. Commun. 194, 1038-1043, 1993
A;Tille: Sequence and functional Characterization of feline interleukin 2.
A;Reference number: JN0698; MUID:93356765
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        interleukin 2 precursor - cat
C;Species: Felis silvestris catus (domestic cat)
C;Date: 03-Feb-1994 #sequence_revision 03-Feb-1994 #text_change 16-Jul-1999
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                                                                                                                                                                                                                                                                                                                                                            A, Gene: GDB:1L2
A; Cross-references: GDB:119344; OMIM:147680
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C;Species: Meriones unguiculatus (Mongolian jird)
C;Species: Meriones unguiculatus (Mongolian jird)
C;Species: Meriones unguiculatus (C;Date: 06-Jan-1995 *text_change 16-Jul-1999
C;Accession: S33509
R;Mai, Z: Klei, T.; Horohov, D.
R;Mai, Description: Cross-species PCR cloning of Jird (Meriones unguiculatus) interleukin A;Reference number: S33509
A;Reference number: S33509
A;Retus: preliminary
A;Molecule type: mRNA
A;Residues: 1155 < Walk.
A;Residues: 1155 < Walk.
A;Residues: 1155 < Walk.
A;Residues: 1155 < Walk.
C;Superfamily: interleukin-2
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C;Species: Equus caballus (domestic horse)
C;Date: 13-Jan-1995 #sequence_revision 13-Jan-1995 #text_change 16-Jul-1999
C;Dates: 13-Jan-1995 #sequence_revision 13-Jan-1995 #text_change 16-Jul-1999
R;Tavernor, A.S.; Butcher, G.W.
Submitted to the EMBL Data Library, November 1992
A;Description: colNA cloning of equine interleukin-2 by polymerase chain reaction.
A;Reference number: S31391
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            caprine interleukin 2.
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C;Date: 06-Jan-1995 #sequence_revision 06-Jan-1995 #text_change 16-Jul-1999
C;Accession: S38662
R;Rimstad, E.
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A;Molecule type: mRNA
A;Residues: 1-149 <TAV>
A;Cross-references: EMBL:X69393; NID:g1076; PIDN:CAA49190.1; PID:g1077
C;Superfamily: interleukin-2
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                               Length 154;
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A;Description: The molecular cloning and expression of
A;Reference number: S38662
A;Accession: S38662
Score 96; DB 2; Len
Pred. No. 6.6e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match 62.6%; Score 92; DB 2; Le. Best Local Similarity 56.7%; Pred. No. 2.5e-06; Matches 17; Conservative 8; Mismatches 5;
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                                                                                                                              4; Mismatches
                                                                                                                                                                                                                                                                                  21 APTSSPAKEAQOYLEQLLLDLQQLLRGINN 50
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                                    Ouery Match 65.3%;
Best Local Similarity 66.7%;
Matches 20; Conservative
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Best Local Similarity 66.7
Matches 20; Conservative
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N. Alternate names: IL-2; T-cell growth factor
C; Species: Sus scrofa domestica (domestic pig)
C; Date: 30-Jun-1992 #sequence_revision 30-Jun-1992 #text_change 16-Jul-1999
R; Goodall, JC.; Emery, D.C.; Bailey, M.; English, L.S.; Hall, L.
A; Reference number: S16241; MUID: 91274360
A; Recession: S16241
A; Residues: 1-154 <GGO>
A; Reference number: S15473
A; Recession: S13473
A; Recession: S13473
A; Recession: S13473
A; Recession: S13473
A; Residues: 1-154 <-LEF>
A; Residues: Life che EMBL: X58428; NID: 92068; PIDN: CAA41330.1; PID: 92069
C; Superfamily: Interleukin-2
C; Superfamily: Interleukin-2
C; Superfamily: Interleukin-2
C; Superfamily: Interleukin-2
C; Reywords: Cytokine; glycoprotein; growth factor; immunoregulation; T-cell
F; 1-20/Domain: signal sequence #status predicted <-MAT>
F; 21-154/Product: Interleukin-2 #status predicted <-MAT>
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                               A; Accession: JN0698
A; Status: nucleic acid sequence not shown
A; Status: nucleic acid
A; Residues: 1-154 <COZ>
A; Cross: references: GB:L19402; NID:g304313; PIDN:AAA02865.1; PID:g304314
C; Superfamily: interleukin-2
C; Reywords: growth factor
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                                                                                                                                                                                                                                                                                                                                                                                           77.6%; Scor.
73.3%; Pred. No. ...
5; Mismatches
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Matches 22; Conservative
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submitted to the EMBL Data Library, December 1989
A;Reference number: $21470
A;Accession: $21470
A;Accession: $21470
A;Accession: $21470
A;Molecule type: DNA
A;Residues: 1-22 <AN2>
A;Residues: 1-22 <AN2>
C;Genetics:
C;Genetics:
A;Cross-references: EMBL:X17201; NID:g452; PIDN:CAA35062.1; PID:g453
C;Genetics:
C;Genetics:
C;Genetics:
C;Cenetics:
C;Cenetic
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R;Ghosh, S.; Palmer, S.M.; Rodrigues, N.R.; Cordell, H.J.; Hearne, C.M.; Cornall, R
Nature Genet. 4, 404-409, 1993
A;Title: Polygenic control of autoimmune diabetes in nonobese diabetic mice.
A;Reference number: S36162; MUID:94004970
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N;Alternate names: IL-2; T-cell growth factor (TCGF)
C;Species: Mus musculus (house mouse)
C;Date: 30-Jun-1987 **Requence_revision 30-Jun-1987 **text_change 21-Jul-2000
C;Accession: A93550; A34490; A94664; I48597; A01850; I84713
R;Fuse, A.; Fujita, T.; Yasumitsu, H.; Kashima, N.; Hasegawa, K.; Taniguchl,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             C;Species: Mus musculus (house mouse)
C;Date: 13-Jan-1995 #sequence_revision 13-Jan-1995 #text_change 16-Jul-1999
C;Accession: S37289; S27205; S36162; S24936
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C;Superfamily: interleukin-2
C;Keywords: cytokine; glycoprotein; growth factor; lymphokine; T-cell
F;1-20/Domain: signal sequence #status predicted <SIG>
F;21-63/Product: interleukin-2 #status predicted <MAT>
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Submitted to the EMBL Data Library, April 1993
A; Reference number: $37289
A; Status: preliminary
A; Moression: $37289
A; Status: preliminary
A; More coule type: mRNA
A; Mesidues: 1-169
A; Cross-references: EMBL:X73040
B; Matesanz, F.; Alcina, A; Pellicer, A.
Biochim. Biophys. Acta 1132, 335-336, 1992
A; Title: A new CDNA sequence for the murine interleukin-2 gene.
A; R; Matesanz, F.; Alcina, A; Pellicer, A.
Biochim. Biophys. Acta 1132, 335-336, 1992
A; Title: A new CDNA sequence for the murine interleukin-2 gene.
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Pred. No. 0.00023;
6; Mismatches 8.
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Pred. No. 0.008;
6; Mismatches
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Local Similarity 53.3%;
es 16; Conservative
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Best Local Similarity 47.2%;
Matches 17; Conservative
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A, Molecule type: mRNA
A; Residues: 1-63 <MATE>
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A; Molecule type: mRNA
A; Residues: 1-50 <GHO>
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S37289
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R;Seow, H.F.; Rothel, J.S.; Radford, A.J.; Wood, P.R.
Nucleic Acids Res. 18, 7175, 1990
Nucleic Acids Res. 18, 7175, 1990
A;Title: The molecular cloning of ovine interleukin 2 gene by the polymerase chain react A;Reference number: S13102; MUID:91088336
A;Accession: S13102
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C; Species: Bos primigenius taurus (cattle)
C; Accession: 145913; S21470; S2071, S
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C;Species: Ovis orientalis aries, Ovis ammon aries (domestic sheep)
C;Species: Ovis orientalis aries, Ovis ammon aries (domestic sheep)
C;Species: Ovis orientalis aries, Ovis ammon aries (domestic sheep)
C;Species: Ovin-1993 #sequence_revision 10-Nov-1995 #text_change 16-Jul-1999
C;Accession: SI488; S13102; S15517
R;Goodall, J.C.; Emery, D.C.; Perry, A.C.F.; English, L.S.; Hall, L.
Nucleic Acids Res. 18, 5883, 1990
A;Title: CDNA cloning of ovine interleukin 2 by PCR.
A;Reference number: S11488; MUID:91016933
            A;Molecule type: mRNA
A;Residues: 1-155 <RIM>
A;Cross-references: EMBL:X76063; NID:g416002; PIDN:CAA53664.1; PID:g416003
C;Superfamily: interleukin.2
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A;Molecule type: mRNA
A;Molecule type: mRNA
A;Residues: 1-155 <CER>
A;Cross-references: GB:M12791; NID:g163204; PIDN:AAA30586.1; PID:g163205
R;Anikeeva, N.N.; Vinogradova, T.V.; Votoshin, O.N.
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A;Molecule type: mRNA
A;Molecule to the EMBL Data Library, April 1991
A;Molecule type: mrs.
A;Molecule type: m
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Pred. No. 0.00023;
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A;Residudes: 21-153 <BUJ>
Cross-references: EMBL:X60148
C;Superfamily: interleukin-2
                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   16; Conservative
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les 16; Conserv
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A; Molecule type: mRNA
A; Residues: 1-155 <GOO>
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Best Local Similarity
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Best Local &
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C;Species: Mus musculus (house mouse)
C;Species: Mus musculus (house mouse)
C;Date: 02-Aug-1996 #sequence_revision 02-Aug-1996 #text_change 16-Jul-1999
C;Accession: 154512
R;Matesanz, F;Alcina, A.; Pellicer, A.
Immunogenetics 38, 300-303, 1993
A;Title: Existence of at least five interleukin-2 molecules in different mouse stra A;Title: Existence of at least five interleukin-2 molecules in different mouse stra A;Reference number: 154512; MUID:93307791
A;Reference number: 154512
A;Reference number: 154512
A;Residues: 1562-RES>
A;Residues: 1-62-RES>
A;Residues: 1-62-RES>
A;Cross references: GB:LO7574; NID:9349513; PIDN:AAA39326.1; PID:9349514
C;Genetics:
A;Genetics:
C;Superfamily: interleukin-2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            interleukin 2 - mouse (fragment)
C:Species: Mus musculus (house mouse)
C:Decies: Mus musculus (house mouse)
C:Decies: 02-Aug-1996 #sequence_revision 02-Aug-1996 #text_change 16-Jul-1999
C:Accession: 168871
Immunogenetics 38, 300-303, 1993
A:Title: Existence of at least five interleukin-2 molecules in different mouse stra. A:Reference number: 154512; MUID:93307791
A:Accession: 168871
A:Accession: 168871
A:Accession: 168871
A:Residues: 1-72 <RES>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     A;Cross-references: GB:L07576; NID:g349517; PIDN:AAA39328.1; PID:g349518
C;Genetics:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 57.5; DB 2;
Pred. No. 0.16;
5; Mismatches 5;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 58.5; DB 2
Pred. No. 0.094;
6; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1 APTSSST-----KKTQLQLEHLLLDLQMILN 26
       56
                                               25 SPTSSSTASSTAEAQQQQQHLEQLLMDLQELLS 57
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35.68;
                                                                                                                                                           interleukin 2 - mouse (fragment)
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Best Local Similarity 35.6
Matches 16; Conservative
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C;Superfamily: interleukin-2
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les 15; Conserv
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R;Yokota, T.; Arai, N.; Lee, F.; Rennick, D.; Mosmann, T.; Arai, K.
R;Yokota, T.; Arai, N.; Lee, F.; Rennick, D.; Mosmann, T.; Arai, K.
R;Yokota, T.; Arai, N.; Lee, F.; Rennick, D.; Mosmann, T.; Arai, K.
A;Title: Use of a cDNA expression vector for isolation of mouse interleukin 2 cDNA clone A;Reference number: A94064; MUID:85113172
A;Accession: A94064
A;Accession: A94064
A;Molecule type: mRNA
A;Residues: 1-169 
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R; Matesanz, F.: Alcina, A.; Pellicer, A.
munoogenetics 38, 300-303, 1993
A; Title: Existence of at least five interleukin-2 molecules in different mouse strains. A; Reference number: 154512; MUID: 93307791
A; Accession: 168870
                                                                                    A; Molecule type: DNA
A; Residues: 1-169 <-ROP
R; Degrave, W.; Simons, G; Devos, R.; Plaetinck, G.; Remaut, E.; Tavernier, J.; Fiers,
Mol. Biol. Rep. 11, 57-61, 1986
A; Title: Cloning and structure of a mouse interleukin-2 chromosomal gene.
A; Reference number: A54490; MUID:86118396
A; Accession: A54490;
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C;Species: Mus spretus (western wild mouse)
C;Date: 02-Aug-1996 #sequence_revision 02-Aug-1996 #text_change 16-Jul-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
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C; Genetics:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  14;
         A;Title: Organization and structure of the mouse interleukin-2 gene. A;Reference number: A93550; MUID:85087940
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       A;Status: preliminary; translated from GB/EMBL/DDBJ
A;Molecule type: DNA
A;Residues: 1-60 <RES>
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Pred. No. 0.06
6; Mismatches
Res. 12, 9323-9331, 1984
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llarity 45.5%;
Conservative
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Best Local Similarity 38.69
Watches 17; Conservative
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C;Superfamily: interleukin-2
                                                                                                                                                                                                                                                  A; Molecule type: DNA
A; Residues: 1-169 <DEG>
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Best Local Similarity
Matches 15; Conserv
  Nucleic Acids
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Interleukin-2 muta Sequence of Interleukin-2 muta Sequence of human Interleukin-2 muta Interleukin-2 muta Interleukin-2 muta Sequence of an interleukin-2 muta Sequence of an interleukin-2 muta Sequence of interles Sequence of interles Sequence of interleukin-2 muta Sequence of interlequence of interlequence of interles Sequence of interlequence of interlequence

Plasmid pilori35-8 Sequence of non-gl Sequence encoded b

Sequence of human Interleukin 2 alal Interleukin 2 alal

Recombinant human

Novel cytokine peptides and antibody for preventing and/or treating undesirable immune reactions e.g. graft rejection and autoimmune

disorders

Sequence of interl Interleukin-2 muta Interleukin-2 muta

Score

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Result

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Alzari

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Human interleukin

score:

Title: Perfect sc Seguence:

OM protein

Run on:

Scoring table:

Searched:

Minimum DB : Maximum DB :

Database

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IL-2; interleukin 2; human; IP130; immunosuppressive; antirheumatic; graft rejection; autoimmune disorder; rheumatoid arthritis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Goldberg M,
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                                           AAR10901
AAR10902
AAR10903
AAR10904
AAP40048
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AMP50857
AMP50859
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AAP60220
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Mazie J;
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N-PSDB; AAZ88838
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  Human IL-2 N-termi
Human Interleukin-
Interleukin-2 ALPH
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Fusion protein con
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Human IL-2 derived
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Human interleukin-
                                                                              October 21, 2002, 09:46:41; Search time 30 Seconds (without alignments) 111.074 Million cell updates/sec
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             GenCore version 5.1.3
Copyright (c) 1993 - 2002 Compugen Ltd.
                                                                                                                                                  1 APTSSSTKKTQLQLEHLLLDLQMILNGINN 30
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Listing first 45 summaries
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                                                           protein search, using
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seq length: 200000000
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1 APTSSSTKKTQLQLEHLLLDLQMILNGINN 30
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Best Local Similarity
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Best Local Similarity
Matches 30; Conserv
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                                                                                                                                                 31 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                           Homo sapiens
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                                          (II) amino acid sequence or its homologous sequence (H) exhibiting the same activity or binding characteristic. The products of the invention have immunosuppressive and antirheumatic activity. (I) or (II) is used for preparing medicament useful to induce IL-2 activity. (A) or a peptide binding to it is used for inhibiting the activity of IL-2R in treating conditions related with undesirable immune reactions like graft rejection, and autoimmune disorders like rheumatoid arthritis. It is also used for immunoaffinity purification of native or recombinant IL-2 peptides. (V) is also used for preparing medicaments in treating patients with deficient IL-2 activity. (III) used to prepare antisense can be used in gene therapy applications. The peptides inhibits or mimics the binding of helix A of IL-2 to a subunit of an IL-2 minics the binding of helix A of IL-2 to a subunit of an IL-2 minics and those produces
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     This invention describes a novel a peptide, IP130 having 31 (I) or 30 (II) amino acid sequence or its homologous sequence (H) exhibiting the same activity or binding characteristic. The products of the invention have immunosuppressive and antirheumatic activity. (I) or (II) is used for preparing medicament useful to induce IL-2 activity. (A) or a peptide
                                                                                                                                                                                                                                                                                                    Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               IL-2; interleukin 2; human; IP130; immunosuppressive; antirheumatic; graft rejection; autoimmune disorder; rheumatoid arthritis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                treating
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Alzari P;
                                                                                                                                                                                                            a specific interaction. This sequence represents an IP130 peptide described in the method of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Novel cytokine peptides and antibody for preventing and/or treat. undesirable immune reactions e.g. graft rejection and autoimmune
                                invention describes a novel a peptide, IP130 having 31 (I)
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                                                                                                                                                                                                                                                                           Length 30;
                                                                                                                                                                                                                                                                                                    Indels
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                                                                                                                                                                                                                                                                          DB 21;
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                                                                                                                                                                                                                                                                                      Pred. No. 1.1e-13;
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                                                                                                                                                                                                                                                                                                    Mismatches
                                                                                                                                                                                                                                                                          100.0%; Score 147; 100.0%; Pred. No. 1
                                                                                                                                                                                                                                                                                                                                        1 APTSSSTKKTQLQLEHLLLDLQMILNGINN 30
                                                                                                                                                                                                                                                                                                                             1 APTSSSTKKTQLQLEHLLLDLQMILNGINN 30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human IL-2 derived peptide IP130 #1
                                                                                                                                                                                                                                                                                                                                                                                                                 AAY51597 standard; Protein; 31 AA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Claim 1; Page 54; 56pp; English.
             Claim 1; Paqe 55; 56pp; English.
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                                                                                                                                                                                                                                                                                                    Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (INSP ) INST PASTEUR
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                                                                                                                                                                                                                                                                                        Best Local Similarity
                                                                                                                                                                                                                                                   30 AA;
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Mazie J;
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binding to it is used for inhibiting the activity of IL-2R in treating conditions related with undesirable immune reactions like graft rejection, and autoimmune disorders like rheumatoid arthritis. It is also used for immunoaffinity purification of native or recombinant IL-2 peptides. (V) is also used for preparing medicaments in treating patients with deficient IL-2 activity. (III) used to prepare antisense can be used in gene therapy applications. The peptides inhibits or mimics the binding of helix A of IL-2 to a subunit of an IL-2 which a specific interaction. This sequence represents an IP130 peptide described in the method of the invention.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Method for expressing heterologous proteins - as fusion protein,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ö
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                                                                                                                                                                                                                                                                                                                                                                                                              Length 31;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Fernandez Maso JR;
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Herrera Martinez
                                                                                                                                                                                                                                                                                                                                                                                                              100.0%; Score 147; DB 21; 100.0%; Pred. No. 1.1e-13;
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100.0%; Pred. No. 2.2e-13;
Live 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human interleukin-2-derived stabiliser peptide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mismatches
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Estrada Garcia MP, Garcia Suarez J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Novoa Perez LI, Machado Lahera JA,
Benitez Fuentes JV, Narciandi Diaz
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Resonances between coding and non-coding regions were measured for the native human IL-2 gene in plasmid pT911 (see AAQ55630) and a synthetic IL-2 gene (AAQ55629) in which alternative, degenerate codons were used in order to introduce additional restriction sites. It was found that the degenerate changes greatly upset the "natural order" between coding and non-coding regions; as a result, the amount of protein expressed by the degenerate gene is likely to be adversely affected. The inventors have proposed an "optimised" IL-2 gene with the aim of increasing the amount of protein expressed by the gene. (N.B. the sequence is also described as
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Targetted hetero-association of recombinant proteins to multi-functional complexes - useful for therapeutic and diagnostic
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Multifunctional protein; targetted hetero-association; protein assembly; antibody engineering; interleukin-2; IL-2; cytokine; tumour; therapy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Length 60;
                                                                                                                                                                                                    Application of optimised gene expression – for scientific, industrial and therapeutic purposes
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  100.0%; Score 147; DB 15; 100.0%; Pred. No. 2.3e-13;
                                                                                                                                                                                                                                                  Disclosure; Fig 28 and Fig 29; 110pp; French.
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                      92FR-0007571.
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                                                                                                                                                                  N-PSDB; AAQ55629, AAQ55630.
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Best Local Similarity
Matches 30; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                   interleukin-6).
                                                                                       (PERE/) PEREZ
                      22-JUN-1992;
                                                      22-JUN-1992;
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                                                                                                                     Perez J;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Immunoprotein MPB 70 encoding sequence may be encorporated into plasmid pT13S with an N-terminal fragment of human IL-2. The plasmid may be used to transform an expression system giving a fusion protein which may be used as a diagnostic agent for bovine tuberculosis infection.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      non-coding region; coding region; resonance; interaction; IL-2; optimization; degenerate sequence; plasmid pT13SNco; pT9-11; gene expression; regulation; recombinant protein production; interleukin-2; interleukin-6.
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                                                                                                                                                                                                                          Bovine tuberculosis; Interleukin-2; IL-2; plasmid pT13S
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         BCG bacteria derived immuno:protein MPB70 - can be used diagnostic agent used to determine bovine tuberculosis.
                                                                                                                                                                                          Human IL-2 N-terminal transcript of plasmid pT13S.
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100.0%; Pred. No. 2.3e-13;
... Mismatches 0;
1 APTSSSTKKTQLQLEHLLLDLQMILNGINN 30
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                                                                                             AAR06838 standard; protein;
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                                                                                                                                                             14-JAN-1991 (first entry)
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N-PSDB; AAQ05976.
                                                                                                                                                                                                                                                            Mycobacterium bovis,
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AAR48245
ID AAR4
                                                               RESULT 4
AAR06838
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Sequences
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                                                                                                                                                                                                                                                                                              AAR78662;
                                                Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Serl M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Region
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Region
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                                                                                                                                                                                                                              RESULT 8
AAR78662
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0
                                                                        the targetted hetero-association of recombinant proteins. The TL-2 is segmented between His79 and Lys80 to form ALPHA (AAR95433) and BETA (AAR95434) segments encoded by MroI-AscI-HinGIII gene cassettes (AAT15267 and AAT15268, respectively). The device combines the low immunogenicity of the plasmatic protein with a preferable effector segmental oysteine bridge (Cys88-Cys105) after assembly. The combination of 1 or more antibody fragments against tumour antigens with additional cytokines, e.g. IL-6 or IL-12, targets the multi-cytokine complex directly to the tumour.
                                                                                                                                                                                                                                                                                                                                 Gaps
                                                           Human interleukin-2 (\text{IL-2}) can be used as an assembly device for the targetted hetero-association of recombinant proteins. The \text{IL-2}
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Pro-insulin is produced by treating recombinant precursor protein with a mercaptan to provide 2-10 SH residues per Cys residue, in presence of a chaotropic agent and in aq. medium of pH 10-11, treating the prod. with 3-50 g hydrophobic adsorber resin per 1 aq. medium of pH 4-7, isolating the adsorbed resin and pro-insulin and desorbing the pro-insulin. This method produces pro-insulin and correctly bonded Cys bridges. Compared with known methods it involves fewer stages (esp. no sulphitolysis or cyanogen bromide cleavage) and overall losses during purification are reduced, i.e. the process is quicker and gives better yields.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Prodn. of pro-insulin with correct di:sulphide bridges - by treating recombinant precursor protein with mercaptan in alkall and in presence of chaotropic agent, then isolation on
                                                                                                                                                                                                                                                                                                                             ö
                                                                                                                                                                                                                                                                                              Length 88;
                                                                                                                                                                                                                                                                                                                               Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Pro-insulin; A-chain; B-chain; C-chain; disulphide; mercaptan; chaotropic agent.
                                                                                                                                                                                                                                                                                              DB 17;
                                                                                                                                                                                                                                                                                           Ouery Match 100.0%; Score 147; DB 17; Best Local Similarity 100.0%; Pred. No. 3.4e-13; Matches 30; Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sabel W;
                                                                                                                                                                                                                                                                                                                                                                            APTSSTRKTQLOLEHLLLDLOMILNGINN 32
                                                                                                                                                                                                                                                                                                                                                             1 APTSSSTKKTQLQLEHLLLDLQMILNGINN 30
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Disclosure; Page 11; 15pp; German.
                             Example 3; Fig 13; 34pp; English,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ×
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAR68899 standard; peptide; 96
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Obermeier
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        92DE-4240420
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       93EP-0118993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human pro-insulin 2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 1994-177718/22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Ludwig J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (FARH ) HOECHST AG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      hydrophobic resin
                                                                                                                                                                                                                                                           88 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         02-MAR-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     25-NOV-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        02-DEC-1992;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      08-JUN-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         EP600372-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAR68899;
                                                                                                                                                                                                                                                               Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gerl M,
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                                                                                                                                                                                                                                                                                                                                                                                                                                            RESULT 7
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Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The present sequence is that of a fusion protein, produced in E.coll which contains an example of a proinsulin molecule corresp.

to the general formula R2-R1-(B2-B29)-Y-x-G1y-(A2-A20)-R3 (II). In formula (II). X - Lys, Arg or a peptide of 2-35 amino acids contg.

Lys or Arg at the N- and C-termini: Y = a natural amino acid; R1 - Phe or a bond; R2 = H, Arg, Lys, a a peptide of 2-45 amino acid; R1 - Phe or a bond; R2 = H, Arg, Lys, a peptide of 2-45 amino acids contg. Arg or Lys at the N- and C-termini: R3 - a natural amino acids acid; (A2-A20) and (B2-B29) are the insulin A- and B-chain sequences from human or other insulin. The proinsulin molecule, released by cyanogen bromide, is reacted with mercaptan at a ratio of 2-10 SH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Proinsulin; post-translational modification; recombinant production; protein folding; conformation.
                                                                                                                                                                                                                ö
of insulin chain A, B and C are given in AAR68895-97 of pro-insulin 1-4 are given in AAR68898-901,
                                                                                                                                                         Length
                                                                                                                                                                                                                   Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 "a peptide of 4 amino acids"
                                                                                                                                                     Query Match 100.0%; Score 147; DB 15; Best Local Similarity 100.0%; Pred. No. 3.8e-13; Matches 30; Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /note- "human insulin B-chain"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /note= "human insulin A-chain"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Fusion protein contg. proinsulin sequence 3.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              76..96
/label- Gly-(A2-A20)-R3
                                                                                                                                                                                                                                                                              1 APTSSSTKKTQLQLEHL,LLDLQMILNGINN 30
                                                                                                                                                                                                                                                                                                                                     31
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              45.74
/label- R1-(B2-B29)-Y
                                                                                                                                                                                                                                                                                                               2 APTSSSTKKTQLQLEHLLLDLQMLLNGINN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Example 2; Page 8; 16pp; German.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAR78662 standard; protein; 96
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               95EP-0101748
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         94DE-4405179
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    03-APR-1996 (first entry)
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/label= 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    [FARH ] HOECHST AG.
                                                                                           96 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        09-FEB-1995;
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Interleukin-2 mutant psI 141 (delta74-78).
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N-PSDB; AAQ10780.
                                                                                                                                  (SERA-) SERAGEN INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (SERA-) SERAGEN INC.
                                                                                                                                                                                                                                                                                                                                                         Ouery Match
Best Local Similarity
Matches 30; Conserv
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                                                                                                                                                                                                                                                                                                                                         128
                                                                                                                                                    Genbauffe FS,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Genbauffe FS,
                                                                                             30-JUL-1990;
                                                                                                                02-AUG-1989;
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                                                         WO9102000-A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           30-JUL-1990;
                                                                            21-FEB-1991
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                10-MAY-1991
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                                        Synthetic.
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                                                                                                                                                                                                                                                                                                                                         Sequence
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                                                                                                      Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                   of IL-2 and and for
residues of mercaptan per Cys residue of proinsulin. The reaction takes place in the presence of a chaotropic auxiliary agent at pH 10-11 and results in proinsulin with correctly linked Cystine bridges. Reaction with trypsin and opt. carboxypeptidase B yields correctly folded insulin. The insulin is isolated by absortion on
                                                                                                                                                                                                                                                          Immunotherapy; lymphokine; interleukin-2; thymocyte mitogenesis
                                                                                                                                                                                                                                         Sequence of interleukin II (IL-2) analogue 341 [Stop121] IL-II.
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                                                                                  Length 96;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Length 120;
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                                                                                                     Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Indels
                                                                                                                                                                                                                                                                                                                                                                                                                    Microbial expression of interleukin II and analogues - manufactured DNA sequences to transform microorganisms
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  gene for the prodn. IL-2 and analogues,
                                                                                100.0%; Score 147; DB 16;
100.0%; Pred. No. 3.8e-13;
Live 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      100.0%; Score 147; DB 6;
100.0%; Pred. No. 4.8e-13;
Live 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                               The inventors claim a manufactured gene for analogues, and for polypeptides of IL-2 and methods for their recombinant production.
                                                                                                                      APTSSSTKKTOLOLEHLLLDLOMILNGINN 30
                                                                                                                              2 APTSSSTKKTQLQLEHLLLDLQMILNGINN 31
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1 APTSSSTKKTQLQLEHLLLDLQMILNGINN 30
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                                                                                                                                                                                   AAP50864 standard; Protein; 120
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Protein; 128
                                                                                                                                                                                                                                                                                                                  84WO-US01252
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83US-0521967
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                                                                                                    Conservative
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                                                                                                                                                                                                                                                                                                                                                                                Souza LM, Stabinsky Y;
                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 1985-062280/10.
                                              a hydrophobic resin
                                                                                          Local Similarity
nes 30; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local Similarity
Matches 30; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAR10906 standard;
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                                                               96 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     120
                                                                                                                                                                                                                                                                                                                                                               (AMGE-) AMGEN
                                                                                                                                                                                                                                                                                                                  09-AUG-1984;
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10-AUG-1983;
                                                                                                                                                                                                                        01-DEC-1991
                                                                                                                                                                                                                                                                                                 28-FEB-1985
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                                                                                                                                                                                                      AAP50864;
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                                                               Sequence
                                                                                Query Match
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                                                                                           Best Loc
Matches
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ID AAR1
XX
AC AAR1
XX
DT 10-M
                                                                                                                                                                  RESULT 9
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In this mutant IL-2 Gln(74), Ser(75), Lys(76), Asn(77) and Phe(78) have been deleted. The mutant IL-2 may be covalently linked to a portion of a toxin molecule ed diphtheria, which is large enough to have cytotoxic activity but small enough not to exhibit general eukaryotic cell binding. The IL-2/toxin hybrid can be used to treat immune disorders involving the IL-2 receptor. The mutant IL-2 on its see also AAQ10775-9, AAQ10781-Q10782.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mutant interleukin -2 molecules binding to interleukin-2 receptor - linked with diphtheria toxin to treat eg interleukin-2 receptor positive malignancies or prevent graft resection
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length 128;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DB 12;
.2e-13;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             100.0%; Score 147; Cilarity 100.0%; Pred. No. 5.2 Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Interleukin-2 mutant psI 143 (delta75-77).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 interleukin-2; IL-2; immunostimulant
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     interleukin-2; IL-2; immunostimulant
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim 2; Page 9; 17pp; English.
                                                                                                                                                                                                             90WO-US04258
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                                                                                                                                                                                                                                                                                                                                                                        Akiyoshi
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Gaps

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Lys(76), Asn(77) and Phe(78) have been deleted. The mutant IL-2 may be covalently linked to a portion of a toxin molecule eg diphtheria, which is large enough to have cytoctoxic activity but small enough not to exhibit general eukaryotic cell binding. The IL-2 Ycoxin hybrid can be used to treat immune disorders involving the IL-2 receptor. The mutant IL-2 on its own can be used as an immunostimulant. See also AAQ10775-Q10780, AAQ10782
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length 130;
Length 129;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mutant interleukin -2 molecules binding to interleukin-2 - linked with diphtheria toxin to treat eg interleukin-2 receptor positive malignancies or prevent graft resection
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Indels
                            Indels
Score 147; DB 12;
Pred. No. 5.2e-13;
Mismatches 0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ó
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mismatches
                                                                                                                                                                                                                                          Interleukin-2 mutant psI 145 (delta76-78).
                                                                                30
                                                       1 APTSSSTKKTQLQLEHLLLDLQMILNGINN 30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1 APTSSSTKKTQLQLEHLLLDLQMILNGINN 30
                                                                                                                                                                                                                                                                     immunostimulant.
                                                                    1 APTSSSTKKTQLQLEHLLLDLQMILNGINN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1 APTSSSTKKTQLQLEHLLLDLQMILNGINN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence of interleukin-2 (IL-2) II.
                                                                                                                                                        Ā
                                                                                                                                                        AAR10907 standard; Protein; 130
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAP30044 standard; peptide; 131
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 4; Page 9; 17pp; English.
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100.0%;
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                                                                                                                                                                                                              (first entry)
                            Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Senbauffe FS, Akiyoshi
                                                                                                                                                                                                                                                                  interleukin-2; IL-2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 1991-073489/10.
                                                                                                                                                                                                                                                                                                                                                                                                                                      (SERA-) SERAGEN INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local Similarity
              Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      130 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           N-PSDB; AAQ10781
                                                                                                                                                                                                                                                                                                                                                                                30-JUL-1990;
                                                                                                                                                                                                                                                                                                                                                                                                           02-AUG-1989;
Query Match
Best Local Simi
Matches 30;
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                                                                                                                                                                                                              10-MAY-1991
                                                                                                                                                                                                                                                                                                                                                      21-FEB-1991
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                                                                                                                                                                                                                                                                                                Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence
                                                                                                                                                                                 AAR10907;
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ID AAP3
XX
AC AAP3
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                                                                                                                       Ser(75), Lys(76) and Asn(77) have been deleted, as has His(79). The mutant IL-2 may be covalently linked to a portion of a toxin molecule eg diphtheria, which is large enough to have cytotoxic activity but small enough not to exhibit general eukaryotic cell binding. The IL-2 /Toxin hybrid can be used to treat immune disorders involving the IL-2 receptor. The mutant IL-2 on its own can be used as an immunostimulant. See also AAQ10775-8, AAQ10780-Q10782.
                                                                                                                                                                                                                                                                                                Gaps
                                      Mutant interleukin -2 molecules binding to interleukin-2 receptor - linked with diphtheria toxin to treat eg interleukin-2 receptor positive malignancies or prevent graft resection
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Lys(76), Asn(77), Phe(78) and His(79) have been deleted. The mutan IL-2 may be covalently linked to a portion of a toxin molecule eg diphtheria, which is large enough to have cytotoxic activity but small enough not to exhibit general eukaryotic cell binding. The IL-2/toxin hybrid can be used to treat immune disorders involving the IL-2 receptor. The mutant IL-2 on its own can be used as an immunostimulant. See also AAQ10775-Q10781.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mutant interleukin -2 molecules binding to interleukin-2 receptor - linked with diphtheria toxin to treat eg interleukin-2 receptor positive malignancies or prevent graft resection
                                                                                                                                                                                                                                                                                              á
                                                                                                                                                                                                                                                                  Length 129;
                                                                                                                                                                                                                                                                                             Indels
                                                                                                                                                                                                                                                                100.0%; Score 147; DB 12; ilarity 100.0%; Pred. No. 5.2e-13; Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Interleukin-2 mutant psI 150 (delta76-79).
                                                                                                                                                                                                                                                                                                                         1 APTSSSTKKTQLQLEHLLLDLQMILNGINN 30
                                                                                                                                                                                                                                                                                                                                          1 APTSSSTKKTQLQLEHLLLDLQMILNGINN 30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    interleukin-2; IL-2; immunostimulant
                                                                                                                                                                                                                                                                                                                                                                                                                       AAR10908 standard; Protein; 129
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 4; Page 9; 17pp; English.
                                                                                              Claim 7; Page 9; 17pp; English
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               89US-0388557
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Akiyoshi
WPI; 1991-073489/10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 1991-073489/10.
N-PSDB; AAQ10782.
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                                                                                                                                                                                                                                                                              Best Local Similarity
Matches 30; Conserv
                                                                                                                                                                                                                                       129 AA;
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             N-PSDB; AAQ10779
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               02-AUG-1989;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             10-MAY-1991
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WO9102000-A
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Synthetic.
                                                                                                                                                                                                                                         Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAR10908;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The inventors claim a cloned IL-2 gene, esp. AAN30031 and fragments and DNA encoding AAN30043-P30045. The gene is used to prod. recombinant IL-2 which is stable at pH9-9 and on heating at 56 degrees C and promotes the growth of monoclonal functional T-cells.
T-cell growth factor; mitogen; antitumour; lymphokine; T-limphokine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence of human interleukin-2 (IL-2) mature protein, sequence II.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     T cell growth factor; lymphokine; lymphocyte reactivity modulator; thymocyte mitogenesis; cytotoxic T cell reactivity.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Cloned gene coded for interleukin-2 poly:peptide - useful in prodn. of stable interleukin-2 prod.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Kashima N;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Length 131;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match 100.0%; Score 147; DB 4; Best Local Similarity 100.0%; Pred. No. 5.3e-13; Matches 30; Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Matsui H,
                                         Escherichia coli x 1776/pIL 2-50A FERM BP-226.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAP50217 standard; Protein; 131 AA.
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                                                                                                                                                                                                                                                                                                                                                                                                                               (AJIN ) AJINOMOTO KK.
(NICA-) JAPAN FOUND CANCER RES.
(GANK-) ZH GAN KENKYUKAI.
                                                                                                                                                                                                                    82JP-0230371.
82JP-0051122
82JP-0215519.
82JP-0229619.
82JP-0234607.
82JP-023407.
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                                                                                                                                                                             82EP-0082509
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     131 AA;
                                                                                                                                                                                                                                                                18-MAY-1982;
15-DEC-1982;
24-DEC-1982;
27-DEC-1982;
03-FEB-1983;
29-DEC-1982;
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                                                                                                                                                                             18-MAY-1982;
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                                                                                                                                 19-OCT-1983
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                                                                                     EP91539-A.
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ID AAP50217

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                                                                                                                                                                                                                                      Prodn. of interleukin-2 polypeptide - by cultivation of Saccharomyces cerevisiae transformed with gene and vector DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length 131;
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                                                                                       Sano K;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   100.0%; Score 147; DB 6;
100.0%; Pred. No. 5.3e-13;
iive 0; Mismatches 0;
                                                                                       Sato T,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1 APTSSSTKKTQLQLEHLLLDLQMILNGINN 30
                                                                                       Hamuro J,
                                                                                                                                                                                                                                                                                                                             Disclosure; Fig 2b; 48pp; English
(NICA-) JAPANESE FOUND CANCER RE.
(GANK-) GAN KENKYUKAI ZH.
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Best Local Similarity 100.0
Matches 30; Conservative
                                                                                    Taniguchi T, Matsul H,
                                                                                                                                              WPI; 1985-124043/21.
N-PSDB; AAN50219.
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